



SEQUENCE LISTING

<110> Yen, Frances
Erickson, Mary Ruth
Fruebis, Joachim
Bihain, Bernard

<120> Methods Of Screening For Compounds That Modulate the
LSR-Leptin Interaction and Their Use in the Prevention
and Treatment of Obesity-Related Diseases

<130> 70.US2.REG

<140> 09/668,558

<141> 2000-09-22

<150> 60/155,506

<151> 1999-09-22

<160> 106

<170> Patent.pm

<210> 1

<211> 23187

<212> DNA

<213> Homo sapiens

<220>

<221> exon

<222> 2001..2356

<223> exon1

<220>

<221> exon

<222> 3540..3884

<223> exon2

<220>

<221> exon

<222> 12163..12282

<223> exon3

<220>

<221> exon

<222> 15144..15200

<223> exon4

<220>

<221> exon

<222> 15765..15911

<223> exon5

<220>

<221> exon

<222> 19579..19752

<223> exon6

```

<220>
<221> exon
<222> 19899..19958
<223> exon7

<220>
<221> exon
<222> 20056..20187
<223> exon8

<220>
<221> exon
<222> 20329..20957
<223> exon9

<220>
<221> exon
<222> 21047..21187
<223> exon10

<220>
<221> polyA_signal
<222> 21168..21173
<223> AATAAA

<220>
<221> misc_feature
<222> 1..2000
<223> potential 5'regulatory region

<220>
<221> misc_feature
<222> 22324..23187
<223> homology with USF2 gene in ref: embl Y07661

<220>
<221> primer_bind
<222> 523..544
<223> upstream amplification primer 17-2

<220>
<221> primer_bind
<222> 1047..1068
<223> downstream amplification primer 17-2 , complement

<220>
<221> primer_bind
<222> 946..963
<223> upstream amplification primer 99-4576

<220>
<221> primer_bind
<222> 1385..1402
<223> downstream amplification primer 99-4576 , complement

<220>
<221> primer_bind
<222> 1096..1115
<223> upstream amplification primer 9-19

```

```

<220>
<221> primer_bind
<222> 1616..1635
<223> downstream amplification primer 9-19 , complement

<220>
<221> primer_bind
<222> 1602..1621
<223> upstream amplification primer 9-20

<220>
<221> primer_bind
<222> 2074..2093
<223> downstream amplification primer 9-20 , complement

<220>
<221> primer_bind
<222> 2036..2053
<223> upstream amplification primer 99-4557

<220>
<221> primer_bind
<222> 2563..2580
<223> downstream amplification primer 99-4557 , complement

<220>
<221> primer_bind
<222> 2084..2102
<223> upstream amplification primer 9-1

<220>
<221> primer_bind
<222> 2483..2500
<223> downstream amplification primer 9-1 , complement

<220>
<221> primer_bind
<222> 2470..2489
<223> upstream amplification primer 9-21 , complement

<220>
<221> primer_bind
<222> 2062..2081
<223> downstream amplification primer 9-21

<220>
<221> primer_bind
<222> 3455..3474
<223> upstream amplification primer 9-3

<220>
<221> primer_bind
<222> 3882..3901
<223> downstream amplification primer 9-3 , complement

<220>
<221> primer_bind
<222> 3775..3792

```

```

<223> upstream amplification primer 99-4558

<220>
<221> primer_bind
<222> 4336..4356
<223> downstream amplification primer 99-4558 , complement

<220>
<221> primer_bind
<222> 4902..4920
<223> upstream amplification primer 99-14419 , complement

<220>
<221> primer_bind
<222> 4444..4463
<223> downstream amplification primer 99-14419

<220>
<221> primer_bind
<222> 6638..6655
<223> upstream amplification primer 99-4577

<220>
<221> primer_bind
<222> 7072..7089
<223> downstream amplification primer 99-4577 , complement

<220>
<221> primer_bind
<222> 7995..8012
<223> upstream amplification primer 99-4559

<220>
<221> primer_bind
<222> 8576..8593
<223> downstream amplification primer 99-4559 , complement

<220>
<221> primer_bind
<222> 9622..9639
<223> upstream amplification primer 99-3148

<220>
<221> primer_bind
<222> 10023..10040
<223> downstream amplification primer 99-3148 , complement

<220>
<221> primer_bind
<222> 9964..9981
<223> upstream amplification primer 99-4560

<220>
<221> primer_bind
<222> 10546..10563
<223> downstream amplification primer 99-4560 , complement

<220>
<221> primer_bind

```

```

<222> 10996..11015
<223> upstream amplification primer 99-14411 , complement

<220>
<221> primer_bind
<222> 10492..10512
<223> downstream amplification primer 99-14411

<220>
<221> primer_bind
<222> 11972..11990
<223> upstream amplification primer 99-4561

<220>
<221> primer_bind
<222> 12481..12501
<223> downstream amplification primer 99-4561 , complement

<220>
<221> primer_bind
<222> 12005..12023
<223> upstream amplification primer 9-4

<220>
<221> primer_bind
<222> 12417..12436
<223> downstream amplification primer 9-4 , complement

<220>
<221> primer_bind
<222> 14102..14119
<223> upstream amplification primer 99-4562

<220>
<221> primer_bind
<222> 14543..14563
<223> downstream amplification primer 99-4562 , complement

<220>
<221> primer_bind
<222> 14431..14448
<223> upstream amplification primer 99-3149

<220>
<221> primer_bind
<222> 14848..14865
<223> downstream amplification primer 99-3149 , complement

<220>
<221> primer_bind
<222> 14748..14767
<223> upstream amplification primer 9-22

<220>
<221> primer_bind
<222> 15198..15218
<223> downstream amplification primer 9-22 , complement

<220>

```

```

<221> primer_bind
<222> 14748..14767
<223> upstream amplification primer 9-24

<220>
<221> primer_bind
<222> 15333..15351
<223> downstream amplification primer 9-24 , complement

<220>
<221> primer_bind
<222> 15002..15019
<223> upstream amplification primer 9-5

<220>
<221> primer_bind
<222> 15333..15351
<223> downstream amplification primer 9-5 , complement

<220>
<221> primer_bind
<222> 15640..15657
<223> upstream amplification primer 9-6

<220>
<221> primer_bind
<222> 16072..16089
<223> downstream amplification primer 9-6 , complement

<220>
<221> primer_bind
<222> 15800..15817
<223> upstream amplification primer 99-4563

<220>
<221> primer_bind
<222> 16179..16199
<223> downstream amplification primer 99-4563 , complement

<220>
<221> primer_bind
<222> 19295..19312
<223> upstream amplification primer 99-3150

<220>
<221> primer_bind
<222> 19729..19746
<223> downstream amplification primer 99-3150 , complement

<220>
<221> primer_bind
<222> 19420..19438
<223> upstream amplification primer 9-7

<220>
<221> primer_bind
<222> 19824..19841
<223> downstream amplification primer 9-7 , complement

```

```

<220>
<221> primer_bind
<222> 19798..19815
<223> upstream amplification primer 9-8

<220>
<221> primer_bind
<222> 20137..20155
<223> downstream amplification primer 9-8 , complement

<220>
<221> primer_bind
<222> 19913..19931
<223> upstream amplification primer 9-9

<220>
<221> primer_bind
<222> 20329..20346
<223> downstream amplification primer 9-9 , complement

<220>
<221> primer_bind
<222> 20139..20157
<223> upstream amplification primer 99-4564

<220>
<221> primer_bind
<222> 20582..20599
<223> downstream amplification primer 99-4564 , complement

<220>
<221> primer_bind
<222> 20238..20256
<223> upstream amplification primer 9-10

<220>
<221> primer_bind
<222> 20645..20662
<223> downstream amplification primer 9-10 , complement

<220>
<221> primer_bind
<222> 20410..20424
<223> upstream amplification primer 9-26

<220>
<221> primer_bind
<222> 20690..20706
<223> downstream amplification primer 9-26 , complement

<220>
<221> primer_bind
<222> 20569..20588
<223> upstream amplification primer 9-23

<220>
<221> primer_bind
<222> 21243..21262
<223> downstream amplification primer 9-23 , complement

```

```

<220>
<221> primer_bind
<222> 20583..20604
<223> upstream amplification primer 9-11

<220>
<221> primer_bind
<222> 21015..21034
<223> downstream amplification primer 9-11 , complement

<220>
<221> primer_bind
<222> 20584..20601
<223> upstream amplification primer 99-15285 , complement

<220>
<221> primer_bind
<222> 20139..20158
<223> downstream amplification primer 99-15285

<220>
<221> primer_bind
<222> 20642..20659
<223> upstream amplification primer 99-15287 , complement

<220>
<221> primer_bind
<222> 20207..20227
<223> downstream amplification primer 99-15287

<220>
<221> primer_bind
<222> 20691..20709
<223> upstream amplification primer 99-15286 , complement

<220>
<221> primer_bind
<222> 20238..20257
<223> downstream amplification primer 99-15286

<220>
<221> primer_bind
<222> 20943..20960
<223> upstream amplification primer 9-2

<220>
<221> primer_bind
<222> 21295..21312
<223> downstream amplification primer 9-2 , complement

<220>
<221> primer_bind
<222> 21013..21031
<223> upstream amplification primer 99-15284 , complement

<220>
<221> primer_bind
<222> 20582..20602

```



```

<223> downstream amplification primer 99-15284

<220>
<221> primer_bind
<222> 21019..21038
<223> upstream amplification primer 99-14407 , complement

<220>
<221> primer_bind
<222> 20571..20589
<223> downstream amplification primer 99-14407

<220>
<221> primer_bind
<222> 21079..21097
<223> upstream amplification primer 99-15283 , complement

<220>
<221> primer_bind
<222> 20638..20655
<223> downstream amplification primer 99-15283

<220>
<221> primer_bind
<222> 21013..21032
<223> upstream amplification primer LSRi9f15s

<220>
<221> primer_bind
<222> 21195..21214
<223> downstream amplification primer LSRi10r14s , complement

<220>
<221> primer_bind
<222> 20354..20372
<223> upstream amplification primer LSRx9f13s

<220>
<221> primer_bind
<222> 20570..20591
<223> upstream amplification primer LSRx9f14s

<220>
<221> primer_bind
<222> 20811..20832
<223> downstream amplification primer LSRx9r13s , complement

<220>
<221> allele
<222> 818
<223> 17-2-297 : polymorphic base G or C

<220>
<221> allele
<222> 1243
<223> 9-19-148 : polymorphic base C or T

<220>
<221> allele

```

<222> 1374
 <223> 9-19-256 : polymorphic base A or G

 <220>
 <221> allele
 <222> 1401
 <223> 9-19-307 : polymorphic base A or T

 <220>
 <221> allele
 <222> 1535
 <223> 9-19-442 : polymorphic base deletion of C

 <220>
 <221> allele
 <222> 1788
 <223> 9-20-187 : polymorphic base A or C

 <220>
 <221> allele
 <222> 2391
 <223> 9-1-308 : polymorphic base G or C

 <220>
 <221> allele
 <222> 3778
 <223> 9-3-324 : polymorphic base C or T

 <220>
 <221> allele
 <222> 4498
 <223> 99-14419-424 : polymorphic base T or G

 <220>
 <221> allele
 <222> 15007
 <223> 9-24-260 : polymorphic base A or G

 <220>
 <221> allele.
 <222> 15233
 <223> 9-24-486 : polymorphic base A or G

 <220>
 <221> allele
 <222> 15826
 <223> 9-6-187 : polymorphic base C or T

 <220>
 <221> allele
 <222> 19567
 <223> 9-7-148 : polymorphic base A or G

 <220>
 <221> allele
 <222> 19744
 <223> 9-7-325 : polymorphic base A or G

 <220>

<221> allele
 <222> 19786
 <223> 9-7-367 : polymorphic base A or C

 <220>
 <221> allele
 <222> 20158
 <223> 9-9-246 : polymorphic base G or C

 <220>
 <221> allele
 <222> 20595
 <223> LSRX9-BM (17-1-240) : polymorphic base deletion of AGG

 <220>
 <221> allele
 <222> 21108
 <223> LSRX10-BM : polymorphic base T or G

 <220>
 <221> allele
 <222> 606
 <223> potential polymorphic base C or T

 <220>
 <221> allele
 <222> 5141
 <223> potential polymorphic base insertion of G

 <220>
 <221> allele
 <222> 7428
 <223> potential polymorphic base insertion of C

 <220>
 <221> allele
 <222> 8394
 <223> potential polymorphic base C or G

 <220>
 <221> allele
 <222> 8704
 <223> potential polymorphic base T or C

 <220>
 <221> allele
 <222> 9028
 <223> potential polymorphic base G or A

 <220>
 <221> allele
 <222> 9950
 <223> potential polymorphic base deletion of GAATGAAA

 <220>
 <221> allele
 <222> 9977
 <223> potential polymorphic base T or C

<220>
 <221> allele
 <222> 10021
 <223> potential polymorphic base A or G

<220>
 <221> allele
 <222> 11878
 <223> potential polymorphic base C or T

<220>
 <221> allele
 <222> 19040
 <223> potential polymorphic base deletion of G

<220>
 <221> allele
 <222> 21363
 <223> potential polymorphic base A or G

<220>
 <221> allele
 <222> 21449
 <223> potential polymorphic base C or T

<220>
 <221> allele
 <222> 21451
 <223> potential polymorphic base G or C

<220>
 <221> allele
 <222> 21454
 <223> potential polymorphic base A or G

<220>
 <221> allele
 <222> 21455
 <223> potential polymorphic base G or A

<220>
 <221> allele
 <222> 21569
 <223> potential polymorphic base T or A

<220>
 <221> allele
 <222> 21683
 <223> potential polymorphic base deletion of C

<220>
 <221> allele
 <222> 21694
 <223> potential polymorphic base insertion of T

<220>
 <221> allele
 <222> 21728
 <223> potential polymorphic base deletion of G

<220>
 <221> misc_binding
 <222> 799..817
 <223> 17-2-297.mis1

<220>
 <221> misc_binding
 <222> 819..837
 <223> complement 17-2-297.mis2

<220>
 <221> misc_binding
 <222> 1224..1242
 <223> 9-19-148.mis1

<220>
 <221> misc_binding
 <222> 1244..1262
 <223> complement 9-19-148.mis2

<220>
 <221> misc_binding
 <222> 1330..1373
 <223> 9-19-256.mis1

<220>
 <221> misc_binding
 <222> 1375..1393
 <223> complement 9-19-256.mis2

<220>
 <221> misc_binding
 <222> 1382..1400
 <223> 9-19-307.mis1

<220>
 <221> misc_binding
 <222> 1402..1420
 <223> complement 9-19-307.mis2

<220>
 <221> misc_binding
 <222> 1516..1534
 <223> 9-19-442.mis1

<220>
 <221> misc_binding
 <222> 1769..1787
 <223> 9-20-187.mis1

<220>
 <221> misc_binding
 <222> 1789..1807
 <223> complement 9-20-187.mis2

<220>
 <221> misc_binding
 <222> 2372..2390

```

<223> 9-1-308.mis1

<220>
<221> misc_binding
<222> 2392..2410
<223> complement 9-1-308.mis2

<220>
<221> misc_binding
<222> 3759..3777
<223> 9-3-324.mis1

<220>
<221> misc_binding
<222> 3779..3797
<223> complement 9-3-324.mis2

<220>
<221> misc_binding
<222> 4979..4997
<223> 99-14419-424.mis2

<220>
<221> misc_binding
<222> 4999..5017
<223> complement 99-14419-424.mis1

<220>
<221> misc_binding
<222> 14988..15006
<223> 9-24-260.mis1

<220>
<221> misc_binding
<222> 15008..15026
<223> complement 9-24-260.mis2

<220>
<221> misc_binding
<222> 15214..15232
<223> 9-24-486.mis1

<220>
<221> misc_binding
<222> 15234..15252
<223> complement 9-24-486.mis2

<220>
<221> misc_binding
<222> 15807..15825
<223> 9-6-187.mis1

<220>
<221> misc_binding
<222> 15827..15845
<223> complement 9-6-187.mis2

<220>
<221> misc_binding

```

```

<222> 19548..19566
<223> 9-7-148.mis1

<220>
<221> misc_binding
<222> 19568..19586
<223> complement 9-7-148.mis2

<220>
<221> misc_binding
<222> 19725..19743
<223> 9-7-325.mis1

<220>
<221> misc_binding
<222> 19745..19763
<223> complement 9-7-325.mis2

<220>
<221> misc_binding
<222> 19767..19785
<223> 9-7-367.mis1

<220>
<221> misc_binding
<222> 19787..19805
<223> complement 9-7-367.mis2

<220>
<221> misc_binding
<222> 20139..20157
<223> 9-9-246.mis1

<220>
<221> misc_binding
<222> 20159..20177
<223> complement 9-9-246.mis2

<220>
<221> misc_binding
<222> 20576..20594
<223> LSRX9-BM.mis1(17-1-240)

<220>
<221> misc_binding
<222> 20596..20614
<223> complement LSRX9-BM.mis2(17-1-240)

<220>
<221> misc_binding
<222> 21089..21107
<223> LSRX10-BM.mis1

<220>
<221> misc_binding
<222> 21109..21127
<223> complement LSRX10-BM.mis2

<220>

```

```

<221> misc_binding
<222> 587..605
<223> potentialsite606.mis1 potential

<220>
<221> misc_binding
<222> 607..625
<223> complement potentialsite606.mis2 potential

<220>
<221> misc_binding
<222> 5122..5140
<223> potentialsite5141.mis1 potential

<220>
<221> misc_binding
<222> 5142..5160
<223> complement potentialsite5141.mis2 potential

<220>
<221> misc_binding
<222> 7409..7427
<223> potentialsite7428.mis1 potential

<220>
<221> misc_binding
<222> 7429..7447
<223> complement potentialsite7428.mis2 potential

<220>
<221> misc_binding
<222> 8375..8393
<223> potentialsite8394.mis1 potential

<220>
<221> misc_binding
<222> 8395..8413
<223> complement potentialsite8394.mis2 potential

<220>
<221> misc_binding
<222> 8685..8703
<223> potentialsite8704.mis1 potential

<220>
<221> misc_binding
<222> 8705..8723
<223> complement potentialsite8704.mis2 potential

<220>
<221> misc_binding
<222> 9009..9027
<223> potentialsite9028.mis1 potential

<220>
<221> misc_binding
<222> 9029..9047
<223> complement potentialsite9028.mis2 potential

```



```

<220>
<221> misc_binding
<222> 9931..9949
<223> potentialsite9950.mis1 potential

<220>
<221> misc_binding
<222> 9951..9969
<223> complement potentialsite9950.mis2 potential

<220>
<221> misc_binding
<222> 9958..9976
<223> potentialsite9977.mis1 potential

<220>
<221> misc_binding
<222> 9978..9996
<223> complement potentialsite9977.mis2 potential

<220>
<221> misc_binding
<222> 10002..10020
<223> potentialsite10021.mis1 potential

<220>
<221> misc_binding
<222> 10022..10040
<223> complement potentialsite10021.mis2 potential

<220>
<221> misc_binding
<222> 11859..11877
<223> potentialsite11878.mis1 potential

<220>
<221> misc_binding
<222> 11879..11897
<223> complement potentialsite11878.mis2 potential

<220>
<221> misc_binding
<222> 19021..19039
<223> potentialsite19040.mis1 potential

<220>
<221> misc_binding
<222> 19041..19059
<223> complement potentialsite19040.mis2 potential

<220>
<221> misc_binding
<222> 21344..21362
<223> potentialsite21363.mis1 potential

<220>
<221> misc_binding
<222> 21364..21382
<223> complement potentialsite21363.mis2 potential

```

```

<220>
<221> misc_binding
<222> 21430..21448
<223> potentialsite21449.mis1 potential

<220>
<221> misc_binding
<222> 21450..21468
<223> complement potentialsite21449.mis2 potential

<220>
<221> misc_binding
<222> 21432..21450
<223> potentialsite21451.mis1 potential

<220>
<221> misc_binding
<222> 21452..21470
<223> complement potentialsite21451.mis2 potential

<220>
<221> misc_binding
<222> 21435..21453
<223> potentialsite21454.mis1 potential

<220>
<221> misc_binding
<222> 21455..21473
<223> complement potentialsite21454.mis2 potential

<220>
<221> misc_binding
<222> 21436..21454
<223> potentialsite21455.mis1 potential

<220>
<221> misc_binding
<222> 21456..21474
<223> complement potentialsite21455.mis2 potential

<220>
<221> misc_binding
<222> 21550..21568
<223> potentialsite21569.mis1 potential

<220>
<221> misc_binding
<222> 21570..21588
<223> complement potentialsite21569.mis2 potential

<220>
<221> misc_binding
<222> 21664..21682
<223> potentialsite21683.mis1 potential

<220>
<221> misc_binding
<222> 21684..21702

```

<223> complement potentialsite21683.mis2 potential

<220>

<221> misc_binding

<222> 21675..21693

<223> potentialsite21694.mis1 potential

<220>

<221> misc_binding

<222> 21695..21713

<223> complement potentialsite21694.mis2 potential

<220>

<221> misc_binding

<222> 21709..21727

<223> potentialsite21728.mis1 potential

<220>

<221> misc_binding

<222> 21729..21747

<223> complement potentialsite21728.mis2 potential

<220>

<221> misc_feature

<222> 22113,22122,22227,22264,22268

<223> n=a, g, c or t

<400> 1

| | | | | | | |
|------------|------------|-------------|-------------|-------------|-------------|------|
| ccataatcaa | gaaaatggat | aataagtttt | ggtgggggatg | tggagaaatt | ggaatcctcc | 60 |
| gtgcattgct | ggtgggaatg | tacaatagtg | cagtcattgg | ggaaaacagt | ttggcagttc | 120 |
| ctcaaaaggt | taaaaataga | actaccaagt | caccacagca | ttccattctt | aggcatatat | 180 |
| tcaaaagaaa | tgaaagcaga | tatttgtaca | ccagtgttca | cagctgcact | atttacaata | 240 |
| gtcaaaaggt | agaaacaacc | taggtccatc | cacaaatgaa | tggataaata | aaacgtagca | 300 |
| tatacatata | atggtacact | agtccgctgt | aaaaagaaat | tttgatctta | ctgcatgcta | 360 |
| catggcttcg | acatactaca | acatggatgg | accttgaaaa | cattattctt | tgtgaaataa | 420 |
| actagacaca | ggacaaatgt | tagacgattc | cacttatatg | aggcacctag | aatgggcaat | 480 |
| ttggtaagca | aagtagaata | gaaattacta | ggggcacagg | tagcagggaa | tggggagtta | 540 |
| ctgtttaatg | gtcacagagt | ttatgttggg | gatgatgaaa | cagtttcggg | gataaagagt | 600 |
| ggtgactggt | acacgacatt | gtgaatatac | ttaatgccac | tgaattttac | acttgaagtg | 660 |
| gttaaagcga | taaatattat | agtttgcata | ttttatcata | aaaatatttt | tttaaacgat | 720 |
| gaagggacgt | gaacgggttg | aaattttata | aaaagtggcc | agggaagggtg | tcactgcaat | 780 |
| ggtgtcctac | aggaggagga | agatcatgtg | gacatctscg | ggaagggtgt | tctggcagag | 840 |
| ggagtagcac | gggcgatggc | tctgaggact | gtgagaagta | tagttggaaa | cagcgaggag | 900 |
| gccaggggtg | ccgaagctga | gtaagccaga | gagagtggga | ggaggtgaga | taagaggggg | 960 |
| aaggtcagtt | tctgctgaga | gtgaggagga | gccacaggag | ggctgtgagc | aggtggacgt | 1020 |
| gatctggctt | gagttttaac | agggccagta | gaacaaagca | cgcctgggta | ccgaaaccag | 1080 |
| ccactggcca | gttggcaacc | tgggggagtc | taacgcgagg | aagcgcccag | ggttccccca | 1140 |
| ggatgcgctt | tccctcgccg | ccacctggag | acagcagagt | cacgcccagc | gctgcgcagg | 1200 |
| ctgatcgccg | cgccgcgccc | ccgccctcgg | tcgcagggtg | ctygttccgg | gaatttctaa | 1260 |
| gcggaaaccg | gtcccaagcc | ccgcgccttc | gctcggcccc | tttaagagcc | agaatttccg | 1320 |
| gagggctgac | ccggggctag | ggatgcccag | gggcgcgaacc | acaagttggg | aacrggtggg | 1380 |
| ggaggtggcg | aaaacttccg | wagtgggaatt | ccaacttttc | ctggccctga | ttccccttgg | 1440 |
| gcatccctga | gggggcagag | cttcccttcc | ggggacttta | gagggttcct | caggtcatct | 1500 |
| aactgggaga | cacaggaggc | ccgaagcgcc | ccccctccac | ccggtccgga | ggaaccccag | 1560 |
| tggaaagtga | gaagtcaggc | gccaccaaca | agcctctccc | agccaggact | ttgcttagac | 1620 |
| tcgctcctcc | cggcagggcg | cacctaggcg | ggtccatcgc | cagccgggga | gagggggttg | 1680 |
| ggcagggagg | gaacaggtgc | gcggcgggac | ccgccctatc | tcaacaggtg | aatcgctcca | 1740 |
| agtgggtctc | ggttgcatgg | atctcgggtgc | gcttggtttg | gccgggagmag | atgggggccc | 1800 |
| gaagggacct | gtggtccgca | ggcgccctcc | cagcgggcca | gtcacttggt | tcggggccctg | 1860 |

| | | | | | | |
|-------------|-------------|-------------|------------|------------|-------------|------|
| ggggacggag | cgcacctggg | tcagcccact | tccggggagg | gaggcagagg | aacccctccc | 1920 |
| cgccgctcac | ccctaagccc | agccctcggc | tcccaccctt | gtgtacctgg | gccgaaccat | 1980 |
| tcaccggagc | gcgcagcggg | tggagtgtgg | ctcggaggac | cgcggcgggt | caagcacctt | 2040 |
| tctcccccat | atctgaaagc | atgccctttg | tccacgtcgt | ttacgctcat | taaaacttcc | 2100 |
| agaatgcaac | aggacggact | tggagttagg | acaaggaacg | gaagtgggaa | ggggaggagc | 2160 |
| gtgcacccct | cctggccttg | gtgcgcgccg | cgccccctaa | ggtacttttg | aagggacgcg | 2220 |
| cgggccagac | gcgcccagac | ggccgcgatg | gcgctgttgg | ccggcgggct | ctccagaggg | 2280 |
| ctgggctccc | acccggccgc | cgcaggccgg | gacgcggctg | tcttcgtgtg | gcttctgctt | 2340 |
| agcacctggg | gcacaggtac | ggggcacggg | gcctctgacg | ctgcggaacg | scggagggaa | 2400 |
| ctgtagaggg | ggatggatgg | agttggaggc | ggcggaagc | gggaagcggg | ggtctcagag | 2460 |
| gctgggacct | tccgatcccc | tgggtcttgg | gcgatctgtt | gcgcgcggga | gtgagaggaa | 2520 |
| ttccccat | gtgcgcgggga | gcgctccccg | cgcccttata | tggaagatag | caggaagtga | 2580 |
| aactccctgg | acgggtgagac | ccggagcggc | agggagaatg | gaactctttg | tggggagggga | 2640 |
| gtggaagacc | gcccgatctc | tgggaaaaga | aaagccggga | tgggacttgg | gcgcaccccg | 2700 |
| ggatttctaa | gttttggagt | aacggggaga | gggcacggga | gggctggatc | agacgcttcc | 2760 |
| tagagggaca | gagacgaagg | aacaatgcct | aggcctcggg | tgggtgtggg | actggggact | 2820 |
| ccccatcccc | cgcacccccc | ccacctcccc | cgggctcggg | attatacgtg | cgtaagagtc | 2880 |
| tgggtgggatg | gattttacgga | cttgaaccgc | acttctgctg | gcaggctttc | acctggatgg | 2940 |
| gatatttggg | tgggtgatgag | gtctttcccc | agacactttt | ggttcagtca | tttgaaatga | 3000 |
| ctttagagta | gggtgaggtg | gtgggaggct | gatggagata | ttgtgggggc | tttagtccct | 3060 |
| ccatggcaaa | gcagttcagg | caaacaactc | catggttttc | cctccaaatt | caaaaaggccc | 3120 |
| cgggtaacct | ggaatccttc | gtagtccggt | ttgaagtggg | gccttgggcg | ctgggggcat | 3180 |
| caacatggcc | atctgggctt | gcctgcccag | gccacacaga | ggccccctgt | tgtgggtgaa | 3240 |
| tggcaaaggg | aagaggggac | tgggtgtggt | cagagggcac | aggctgggaa | gagggatggc | 3300 |
| gggcgagtc | aaggaaaactg | gccgtgtcac | cgtgcacctg | ccacttcagc | cccacgggtc | 3360 |
| tataaaatgg | gcatgattat | cgtggctacc | tcactgggtc | tggcaattaa | ggaacaatgt | 3420 |
| gtgccaggca | ctctgtaaac | cacatacttg | cgagtgtcaa | gctggtgaca | ggtggcgctc | 3480 |
| ctgttgaagc | acctccctga | gtcacagca | acccttgcgt | tctctcctct | tgccctcagc | 3540 |
| tcctgccagg | gccatccagg | tgaccgtgtc | caacccctac | cacgtggtga | tcctcttcca | 3600 |
| gcctgtgacc | ctgccctgta | cctaccagat | gacctcgacc | cccacgcaac | ccatcgctcat | 3660 |
| ctggaagtac | aagtctttct | gccgggaccg | catcgccgat | gccttctccc | cggccagcgt | 3720 |
| cgacaaccag | ctcaatgccc | agctggcagc | cgggaaccca | ggctacaacc | cctacgtyga | 3780 |
| gtgccaggac | agcgtgcgca | ccgtcagggt | cgtggccacc | aagcagggca | acgctgtgac | 3840 |
| cctgggagat | tactaccagg | gccggaggat | taccatcacc | ggaagtatgt | tgggcagggc | 3900 |
| agggggatga | ggctgggctt | gcccgggtgg | tgggactggc | gtccttgtgc | gggacctgga | 3960 |
| gtccccatct | gaaagctctt | gagtgccagt | gtctgaaagg | accattgaag | ggagcaattc | 4020 |
| tttttttttt | tttttttgaa | gatggagtct | tgtctgggac | tccaggctgg | agtgcagtgg | 4080 |
| tgcatctca | gctcactgca | acctccacct | cccaggttca | agcaattctc | ttgcctcagc | 4140 |
| ctcccagata | gctgggactc | caggtgcgtg | ccaccacgcc | cagttaattt | ttgtattttt | 4200 |
| agtagagatg | gggtttcacc | atgttggcca | ggctgggtct | aaactcctga | cctcaaataga | 4260 |
| tctgcccgc | ttggcctcgc | aaagtgtctg | gagacaccat | accagacctc | aagggaagca | 4320 |
| ttctattcta | ctattcttcc | ttctgtctaat | ccttccattc | tttaatttaa | taacgaagat | 4380 |
| tttttgagta | cctgtcatat | accaggtgct | gttctgggcc | ctgggaatac | agctgttaac | 4440 |
| aaaatcatca | aaccacttcc | ctcgtggagc | ccacattgca | gtgagagaga | caaacackac | 4500 |
| acacactctc | aagtccttga | agataaagaa | aactgggtaa | cggagagaag | aggccagggg | 4560 |
| ttgttctata | atcattaata | acacgagcag | taagaagtaa | aatttatcta | agtaacaact | 4620 |
| tataaagggt | ctactgtgtg | ctaagctctc | atccagggtc | ccaaggatta | actcagacca | 4680 |
| cacagtaatt | gaatagattc | tatcattgtc | atcttacaga | ggcccagaga | gagaaagtga | 4740 |
| cttgccctagt | gtcatagctg | gtaacggggc | tgggattcta | actcagccac | tttgggtcta | 4800 |
| gtggccaagc | tcctaataccc | tttgcttgcc | taggggtggc | cgcagaggac | tcacagagga | 4860 |
| gatggcagga | gtgaactgca | ggggcaagag | agcttaattg | agaaagcctg | tgacatgcca | 4920 |
| ggaactgcac | acataattctc | ccattgagtc | ctctcctcta | ccctcctgac | agctgaggca | 4980 |
| cagagagggt | accttgttca | aatgggtgca | taggaagtca | aagtctggag | ctggggtttg | 5040 |
| aaccaggca | gccctgagaa | ccttgttctt | tttttttgag | acggagtctc | gctctgtcgc | 5100 |
| ccaggctgga | gtgcagtggc | gggatctcgg | ctcactgcaa | gctccgcctc | ccgggttcac | 5160 |
| gccattctcc | tgccctcagcc | tcccaagtag | ctgggactac | aggcgcccg | cactacgcct | 5220 |
| ggctaatttt | ttgtattttt | agtagagacg | gggtttcacc | gttttagccg | ggatgggtct | 5280 |
| gatctcctga | cctcgtgatc | cgcccgccct | ggcctcccaa | agtgtctggg | ttacaggcgt | 5340 |
| gagccaccgc | gcccggcccc | ttgttcttaa | ctgtaatgct | gcctcctgat | aggatgtgcc | 5400 |

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|-------------|------|
| tggtgggact | aagtaagggg | cagtcattca | ttcattcatt | tggtatttat | caagcatcga | 5460 |
| ctatgtgtcg | ttgggtgctg | ggatagaggt | gattgggatg | gctgaagttt | ctgtcgtcaa | 5520 |
| ggagatgaca | ttctgggtga | gtgagactgg | cagtaaataa | gcagataaag | aaagagtatg | 5580 |
| agaatttcaa | agtctgggca | cgggtggctca | cgtctgtaat | ctcagcactt | tgggaggcca | 5640 |
| aggtgggtgg | atcacctgag | gtcaggagtt | ccagaccagc | ctggccaaca | tggtgaaacc | 5700 |
| ccgtctctac | taaaaataca | aagattagcc | aggcatgggtg | gcacatgcct | gtaatcccag | 5760 |
| ctactcagga | ggctgaggca | tgagaatcgc | ttgaacccag | gaggcagagg | ttgcagtggg | 5820 |
| ctgagatcgc | accactgtac | tgcagtctgg | gcgacagagt | gagactctgt | ctcaaaaaaa | 5880 |
| aaaaaaaaaa | aaaagactcc | gtcaaggtat | aagaatgtca | gagagtacta | agtgttgcaa | 5940 |
| agaaaataac | accaggctgg | gtgcattggc | tcatgcctgt | aaatttcagc | actttggggg | 6000 |
| gccaaggcag | gaggatcact | tgagcctagg | agtttgagac | cagcctggac | aacaaaatga | 6060 |
| gaccccatgt | ctacaaaaat | tttaaaaaat | taaaaattag | ctgggcatgg | tggcatgtgc | 6120 |
| ctgtggtccc | ggctgctcag | gaggctgagg | tgggaggatt | gcttgggctt | gagaggtcaa | 6180 |
| ggcttcagtg | agtcatgatc | gtgccactgc | attccagcct | gggtgacaga | gtgagaccct | 6240 |
| gtcttgaaat | gaaaagaaaa | taggctgggc | gcagtggctc | acacctgtaa | tcccagcact | 6300 |
| ttgggaggcc | gagggtgggtg | gatcacctga | ggtcaggaga | tcgagaccag | cctggccaac | 6360 |
| atgggtgaaat | cccattctcta | ctaaaaatac | aaaatttagc | cgggctgtgt | ggtgggcgcc | 6420 |
| tgtaatccca | gctactcggg | aggctgaggc | aggagaatcg | cttgaacctg | ggaggcgaag | 6480 |
| gttgcgggtg | gccaagattg | cgccactgca | ctctagcctg | ggaaacagtg | agactccgtc | 6540 |
| ttaaaaaaa | aagaaaaaag | aaaatagcac | tgggtgatgt | gctacatgga | atgacttggg | 6600 |
| ctgtgaatat | gatttgagga | gggcctgggc | ctgggcctta | cagaacctag | aaggcagaga | 6660 |
| ggaaggggag | gggcagggtg | ccagggatga | aggctcacgt | acctcatgtc | ttagtgtgtg | 6720 |
| ttcactgtct | taaacaagaa | tttaaagtgt | ggcatggggc | agagcgggga | agggagcatc | 6780 |
| cctttgcaga | ccccaaaga | ccaggaactg | gagcacattc | tgctagagga | tcgatgggaa | 6840 |
| gcagggttcc | aggggctgag | cctatgtcag | tccgttttca | gaggaggcac | caggcttgct | 6900 |
| tgccctgaat | ttctgtgggc | agctcagcca | tgagcatcct | actgttattg | aggctcacagg | 6960 |
| gctgcttagg | ccccctctc | tctaaccag | ggattgtgcc | tgccctggacc | aggcgtgact | 7020 |
| gctaagcttc | tgccaggaca | agccaaatac | tgagggtgct | tcctctgctg | gacgcaaaa | 7080 |
| tccaggatga | ccccccaggc | tctgtctcgg | ggaagggggc | ctgcatgctc | caggggcctc | 7140 |
| acaggcctgg | gtctttcaaa | ccacccccac | ctgggcctgt | gtttgatcaa | ggccctgagt | 7200 |
| gtaaacatcc | attgtgtgtg | tcctttcagg | aaatcccata | gccataggag | cttcctctgt | 7260 |
| ttcagctttg | aggatgggga | aaagtggact | ccccgtgggt | ttcctagggt | caccactgtg | 7320 |
| gctgggggtt | ttctgtgtgt | gttgtttttt | ttctgttgcc | caggctggag | tgcatgggtg | 7380 |
| caatctcagc | tcactgcaac | ctctgcctcg | caagttcaag | tgattctccc | gcctcagcct | 7440 |
| cctgagtagc | tgggattaca | ggtgcacacc | accacacctg | gctaattttt | gtatcttttt | 7500 |
| ggtagagatg | ggatttcgcc | atgttggcca | ggctgggtct | aaactcctga | cctcagggtga | 7560 |
| tctgcctgcc | ttggcctccc | aaagtctctg | gattacagat | gtgagccacc | atgcccggcc | 7620 |
| tatcctgggt | tcaaaagtga | aaatagtctt | ggataaggta | gaaggctgtc | cactccaggc | 7680 |
| atccctccgg | tccggtggct | cattccctgc | tttgtccttc | catgcttttg | gtgatggacc | 7740 |
| agcacctgga | caggaggccc | tgttccacct | cctcgggctc | cttgggggtc | aagtgcccc | 7800 |
| acctccagct | gcactgcagc | agagagccca | tgggacctct | gaaatcatga | aggtcacctt | 7860 |
| tgcggtgtat | aaagaaggaa | ccagaggttg | gagatgtgga | ggaggcctgg | ctgctgttcc | 7920 |
| cactggagac | ctggcatctt | ctccccgacc | taaaacaatg | aaagcagtg | tcagcccgga | 7980 |
| tgagatcacg | gccagcccaa | gaccaggaac | agggtacgcc | ctgcaggaag | aagggtgtgc | 8040 |
| cagaccttag | gatggatcaa | aagaagccgg | aaaactatat | tttttgtgag | ttttgaaaat | 8100 |
| gtcagacagg | tcaaacaaaa | cacagtggag | tccagcctcg | gcctacaaga | tgccagattt | 8160 |
| caacccttgg | cctatatgat | ctgttttgcca | tggcaggcgg | ttcctgtcca | cctcttttgt | 8220 |
| ttatagcagg | gaccagctct | tgagctccag | tggtgaagag | gcacgggtcag | ggtctgatct | 8280 |
| gaagacactg | gtggctcatg | cctgtaatcc | cagcacttca | ggaggccgag | gcaggaggat | 8340 |
| tgcttgagga | caggagctgg | gagaccagcc | tgggcaacac | agtgaagacc | agacactaca | 8400 |
| aaaaaataaa | tttagcgggg | catgatggca | caccctgcta | ctctggagat | gggaagattg | 8460 |
| cttgagcccta | ggaggttcgaa | gctgcagtga | cccactgatc | caccactgca | ctccagcctg | 8520 |
| ggcgaccaag | ctaggccctc | tcaaaaaaga | tacagggtgga | aaaatgatgg | acgaagaggg | 8580 |
| cattgtggca | aacctggggg | tttaggagaa | cctagtttgg | aattctatga | ggattcaatg | 8640 |
| aaagaatgtg | tgtagagggg | cccagcacat | agtaagagct | caataaacgg | tgggggctag | 8700 |
| gggtgggtgg | tcatgcctgt | aatcccagca | ctttggggag | ctgaggcagg | tggatcactt | 8760 |
| gagccctgga | gttcaagatc | aacctggaca | acaaagcaag | atcccatctc | aaaattaaaa | 8820 |
| aacaacacca | acaacaaaaa | aacagtggct | tagatgcctg | atcattaggg | taagtcgtgt | 8880 |
| cctcaacccc | ttcacatctg | ctctgaaggt | caccatatcc | ggaagccttc | cctggcctcc | 8940 |

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|-------------|-------|
| ttgttttaaaa | tggcacagcc | cccactccac | gcctggcact | ctctgctgtc | cctgattcgt | 9000 |
| tttctccata | cagcttatct | ttgtctggta | tgtgacatag | ttaacatttt | atattttgtct | 9060 |
| ttcttttecta | gttagaatct | gaactctaga | agggcaaggg | caaggattta | taactcaaa | 9120 |
| attccgggct | taggcctctt | ttatattctt | gatttttgagg | ttaatttaaga | gttcaggcct | 9180 |
| agcgaggtgg | ctcatgcctg | gaatcccagc | actttgggag | gcccaggcgg | gcagatcact | 9240 |
| tgaggtcagg | agttccagac | ctgcctggcc | aacacagtga | aaaacctgtc | tctactaaaa | 9300 |
| atacaaaaat | tagccagtta | tgttggcagg | cgctataat | cccagctact | caagaggctg | 9360 |
| aggcaggaga | atcgcttgaa | cccaggaggc | agaggctgca | gtgagccaag | atcggtgccac | 9420 |
| tgcactccag | cctgggcaac | agagcgagac | tccatctcaa | aaaaaaaaaa | aaaatttaaga | 9480 |
| gctcaaagag | tttgttttca | taggcagcag | aatgagaaaa | gtttacaaaa | tagtttaaat | 9540 |
| gacaataaag | tcattataga | ttaacataaa | taaaatacct | tttatgaaaa | aaataatcat | 9600 |
| tttctgaaat | cagacaaaac | attgtgaatg | agaaggtggc | atggttttat | ttttttgcaa | 9660 |
| gtctccgaag | cctggctgga | tagaagagcc | tggcttctca | gagctgcttc | agctgtgtgt | 9720 |
| gatatctatt | gtatgtcacg | tagcctctgg | aaaactccac | agttagtatt | gttgggaaaa | 9780 |
| taactttgac | ctcaggatct | cctgaaaacg | tcttggggaa | ccccagggtc | tagaggctgc | 9840 |
| agttttgagaa | ctgttgctgt | ggatatccag | gtgtctcaaa | tactgcctag | aacatagggtg | 9900 |
| gtactcagta | attattgttg | aaggatgaat | gaatgaatga | atgaatgaat | gaaagaaaga | 9960 |
| aatgtgtctt | tgaatctagc | catgtgcccc | gaatgatgag | acagatgaca | aaagctaagg | 10020 |
| gacttttagca | tgaggagagg | gggttcgttt | cctttttttt | cttttttttt | tgagatggag | 10080 |
| tctcactcta | ctgcccaggc | tagagtgcag | tgggtgcaatc | tcagctcact | gcaatctctg | 10140 |
| cctcctgagt | tcaagcaatt | ctcctgcctc | agcctccagg | gtagctggga | ctacagggtgc | 10200 |
| gtgccaccat | gcctagctaa | ttttttacat | ttttggtaga | gatgggggtt | taccatgttg | 10260 |
| gccgggctgg | tctggaactc | ctgacctcaa | gtgatccacc | tgccctcagc | tcccaaagtg | 10320 |
| ttaggattac | aggtgtgagc | caccatgtcc | ggccaagagg | gtgttcattt | ctgctccttg | 10380 |
| ccaggatttg | tgtcaggcac | tggggaccca | gcagtggctg | agacagacag | ggctctgcct | 10440 |
| cacggagccc | acattttcac | caggcaaagg | atggctggcc | cctaagctgg | gagataagac | 10500 |
| ttcagcagtt | gggtggggga | gccgtgggag | aagcccagcc | cacaggggga | cagtgc aaat | 10560 |
| ctagaaccaa | ggcgatggca | gggggtgaggc | tggcacggta | gctagagacc | acgtcgtgcc | 10620 |
| aagggccttg | gggaccatgg | gactatggga | ccttagggaa | ggcgtctgga | atgctgtagc | 10680 |
| cagacactgt | tgcaaggagg | atttttctgt | agacatgagg | ccttccttat | gaagaaagca | 10740 |
| agggttcttt | cattcctggg | ggtgccagg | gctgtggact | gcagcacgcg | tggttgctgc | 10800 |
| cgtcacagag | ctgtcatgca | ggagggcagc | gcgtccttgg | gaagggtggca | ggcaggtcag | 10860 |
| gctaggagga | aagaggccgg | gaagctgagg | gcatttcctg | cccagatgac | ccaatgtagc | 10920 |
| ctacttctgt | ccccagtggc | ttaaggcaga | ggtgcctgg | aggtgccctg | gtccaccct | 10980 |
| ggtgaaaggc | tgaaggatt | taattagtgc | ctgagaagca | gagaggaaac | aggatgtgcc | 11040 |
| aaaacacttt | gatggatgg | agagttaaca | ggctccttgc | ctgcagctgc | ttcagacaag | 11100 |
| agcgtcccca | agccctgggc | ctgacctgga | atgtggggat | ggaaggggag | ggggagggaac | 11160 |
| caaggcactg | ggagggttaag | tctctctctc | ccacatagac | acacccactc | ccttatgggtg | 11220 |
| cctgggcac | tcctggtacc | tagaatctgg | cctgtttatc | tcacacccca | tccttgggg | 11280 |
| ctacactagg | ccctgtgggt | ggcagttcac | atcaggggag | ttctgacttt | ggctctgaga | 11340 |
| ggtggttcag | agatggctgt | aagttgagaa | gcacagactg | ctgggtgtgg | tgggttcacgc | 11400 |
| ctgtaatccc | agcactttgg | gaggctgagg | tgggggtgga | tcacctgagg | tctggagtgc | 11460 |
| aaaaccaact | tggtcaacat | ggcgaaactc | catctctact | aaaaatgcaa | aaattagcca | 11520 |
| ggtgtggtgg | caggtgccta | taatcccagc | tacatgggag | gctgaggcag | gagaatcgct | 11580 |
| tgaatctggg | aggcgaagat | tgtagtgagc | cgagattagt | tcgcaccatt | gcatgccagc | 11640 |
| ctgggcaaca | agagtgaaac | tccgattcaa | acaaacaaaa | aaaaaaagct | gggcatgggtg | 11700 |
| gagtgcctgt | agtcctaact | actcaggtgg | gaggattgct | tgagtccagg | aggttgaagt | 11760 |
| tgcagtgggc | tataattaca | ccactgcact | ccagccagg | ccacagagt | agaccctgtc | 11820 |
| tctaaagaaa | gaaaaaaaaa | aacaacctca | ggctccgagg | gcaccattac | tgctctacac | 11880 |
| tgaagagctg | tgcagctttt | ccagacccga | aatgtcatcc | acaaaacaga | agtataatg | 11940 |
| gtcctgcctc | acagacttct | tgcagtagtc | caggtgttta | gaacgggggtg | taaaaggccg | 12000 |
| tgtgcctctg | gtaggaatct | ttgcataatg | atttgatcat | ctgcagcctg | cccagccac | 12060 |
| tgttgcctcc | cctctgggtg | tgtctgggaag | gggtcttttg | ccctccagg | gttaggtgcc | 12120 |
| ccagcctcca | aggtgccttc | acgccttttc | atcccagactc | agatgctgac | ctgacctttg | 12180 |
| accagacggc | gtggggggac | agtgggtgtg | attactgctc | cgtgggtctca | gcccaggacc | 12240 |
| tccaggggaa | caatgaggcc | tacgcagagc | tcatcgctct | tgggtgagtgg | gcctgggaag | 12300 |
| ggggaggcat | ggcccttct | tttgtccgct | tctgttctgt | ctgcctctcc | ctgtgtccgc | 12360 |
| cctctgcct | ccagcttacc | ctctgggctc | tgtgcctgc | tctgtctctc | cccaggctct | 12420 |
| gccagtcact | taggtctccc | tgtgcctgc | acccaggca | gggaccactg | gcccacagt | 12480 |

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|-------------|-------|
| cctccaatca | cccaagccaa | actaagagaa | gagtggagac | aattggagac | tctgcctttt | 12540 |
| caaagtctca | tttttaaaaa | aatccagac | ttgggggtccg | ggtgcggtag | ttcatgcctg | 12600 |
| taatcccagc | acttttgggag | gccgaggcgg | gtggatcact | tgaggccagg | agttcgagac | 12660 |
| tagcctggcc | aacgtggcaa | aatcccgtct | ctataaaaaa | tataaaagcc | aggcgtgggtg | 12720 |
| gtgcacatgc | ctgtaatccc | agttactcag | aaggctgagg | catgaggatt | gcttgaacct | 12780 |
| gggaggcaga | ggatgcagta | agccaagatc | aagccactgc | actccagcct | gggcgacaga | 12840 |
| gtgagactct | gtccaaaaaa | aaaaaaaaatc | cagacgtggt | cagagtccat | gggcagtga | 12900 |
| tgaggacagt | tgatgggtgtg | caaaatcgac | ccacctcttg | ctacatcccc | aaggcctcat | 12960 |
| ctcacccgag | tccctcgcca | aagcacagcg | gttttgccgt | gtgccctgct | gggatggcgc | 13020 |
| tgcattggcac | acacactgtg | taagtttgag | tgcagctgaa | acgaagccga | ttccagacac | 13080 |
| ccaggggcag | ggcgggggtgt | ccgtgtggct | gggaggcctc | cttgtgttag | ggggatggtg | 13140 |
| ccatcgccca | gggtgccctgc | tgtaaagccaa | cacatggagt | cttgtatgac | atgtgctctg | 13200 |
| catgagtgat | gccgctggggc | tgtacactgc | catcttcaca | tgtgtgaatg | agcacgtgac | 13260 |
| tgggggggtac | ttgggctgca | agacagagtt | catgtgtggg | ggatggaaca | cgtgcaccag | 13320 |
| tgaccaggga | acctctgcct | gttcttcggg | aaaatgcacc | atgtgcatca | gcagttccca | 13380 |
| aaattagtct | ccaggtctat | ttacactcta | aaacattatc | gagggtctcc | aagagctttt | 13440 |
| gtttgtttct | gtgggttttta | tgtctatctg | ttgcttaaca | tattaggaat | taaaatgggg | 13500 |
| agatttttct | tttttttttt | ttttttttga | gatggagtct | cgttctgtcg | cccaggctgg | 13560 |
| agtgcagtgg | ctcgatctcg | gctcactgca | agcttcacct | cctgggttca | cgccattctc | 13620 |
| ctgcctcagc | ctcccaagta | gctgggacta | caggcaccgc | ccaccacacc | cggttaattt | 13680 |
| tttttgtatt | tttagtagag | actgggtttc | accatgttag | ccaggatggt | ctcgatctcc | 13740 |
| tgacctcgtg | atccaccac | ctgggcctcc | caaagtgtcg | ggattacagg | catgagccac | 13800 |
| tgccggcct | taaaatgggg | agatttttca | agcccaagat | acacaaggaa | gactgggcaa | 13860 |
| catggcaaga | ccctgactct | acaaaaaatt | ttaaaattaa | ccaggcatgg | tggcatgcac | 13920 |
| ctgtgagccc | agcttcttgg | gaggctgagg | caggagtatc | gcttgacacc | aggagggtcaa | 13980 |
| ggctgcagtg | agccatgact | atgctactgc | actctagcat | gagtgcacaga | gaccctggct | 14040 |
| caagaaacac | aaacacacac | acacacacac | acacgcata | agtccattag | gcatcagggc | 14100 |
| gatgatggca | tcaggggagcc | tgggaaactc | tactggacat | tcatgggaga | acaagtga | 14160 |
| aaggcaaata | acatcttagt | gttattctaa | aatttcttct | tttggccttg | tggacaggac | 14220 |
| cacgctttga | gagctgtgac | tgacatgcct | ctgtcctgtt | gcgagggcct | atagtgccaa | 14280 |
| gtgcatgagc | tctggggagg | gcttcgtggg | tgcagagctg | ggcctgtgga | ggccctcag | 14340 |
| acacaacact | ggtgggggctc | agagctccag | gggcactcga | gggaagacaa | gaaccggctc | 14400 |
| tgagatgcgt | gaatgtgaca | gtgcatgagt | agagatggag | accttgtggg | tcccagaacc | 14460 |
| aggactgcac | atgactttca | tatgtgggta | tttttgccct | catgggtccc | ttcctgtttt | 14520 |
| aaaaaaaatg | tgtgattatg | ttgtcacaaa | gagtttattc | ctgtatatgt | tgttaatttg | 14580 |
| tgttcagatt | tgtaaagtaa | aattaaacca | tttcagccag | gtgtggtgac | acatgcctgt | 14640 |
| agccctagct | acttacccca | gaggctgagg | tgggaggatc | gcctgagccc | acgaggttga | 14700 |
| agctgcagtg | agccatgac | acacccctgc | actccagact | gggcgacaga | gctgagatcc | 14760 |
| tatttcgtgg | gccctaggtc | cctgtgcctg | ctggaacagg | acatccctat | caccgtgggt | 14820 |
| ggagcccttt | ggggtgctaa | gacctatgaa | tgagggaac | ttagggtgcc | caagctgagg | 14880 |
| tagagccctc | agaacccct | gggatttgta | ttggagccct | cgtggcataa | cacagggtga | 14940 |
| ttatgcaatg | ggagtttctt | acctataagc | accacatgt | gggcgggtgg | agggtaggag | 15000 |
| ccatgcrcta | gggtctcagc | ccccagcccc | ttcccgttc | agggcacacc | ttgcacttgg | 15060 |
| ccagcctgga | gctgggcttt | cgggggtggc | acagcctggg | ctggctctgg | ccagcataat | 15120 |
| ctgtttctct | tttgtccctc | caggaggagc | ctcaggggtg | gctgagctct | tacctggttt | 15180 |
| tcaggcgggg | cccatagaag | gtacgggggg | tggatcctga | gttgggcttc | tcrggagctc | 15240 |
| ccatacatca | cctactgctt | ctgactctag | ttagtatccc | cttccccact | aaaccctgct | 15300 |
| cactgtggac | ccctcactaa | cctggcctga | ctgtggctct | gaggcatcta | gtggtctggc | 15360 |
| gctgggccta | ggctaggctg | ggctgaggag | agcctggggt | gcaggccagg | gctctgtgac | 15420 |
| tggcacctgc | ggtgctcttg | agggtgtggc | gtcctggcag | ctggctctct | ctttgggtctg | 15480 |
| ggggctgcag | tctgtctccc | tctgtgcagg | ctgcctcggt | ttctgccttg | tgttttttgc | 15540 |
| acctggggga | gggcgtaac | tggggaatgg | ccgggatggt | agaatgggga | gtgtgctgtg | 15600 |
| ccagcctct | ggcacaaaaa | atccagccag | ggctgcaggt | tccttggtga | gctttgcaaa | 15660 |
| tcgtccccga | cctcagtgtc | ggctccgcac | catgtacccc | tgtgtgccc | ttagccctgt | 15720 |
| tccctcccag | gcctccgggc | tcagggcctg | ttgtctttct | gcagactggc | tcttcgtggg | 15780 |
| tgtgggtatgc | ctggctgcct | tctcatctt | cctcctcctg | ggcatygtct | ggtgccagtg | 15840 |
| ctgcccgcac | acttgctgct | gctacgtcag | gtgcccctgc | tgcccagaca | agtgtgtctg | 15900 |
| ccccgaggcc | cgtaagtgtc | ccgctcatgg | ccaccctggt | ttgggcaaca | tctgcatcc | 15960 |
| aagggaagga | ggtggccatc | cacctgcccc | caggacagtg | gcgttggtct | ggaggggtgtg | 16020 |

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|-------------|-------|
| aatttagcca | gtggggagaa | agtaggctga | ggagggtctg | ctgttttagat | tgctgtttac | 16080 |
| ttcctccaac | tttttagttta | tttttatatta | tggtgttctt | ttcttttgta | agtataatcc | 16140 |
| atacacatgg | taaaaatgtc | caacagtcaca | agatactagt | cacatggaag | taaagccctc | 16200 |
| taaaaaaacc | aaatcttggc | taggcgcagt | gattacgcct | gtaatcccag | cactttggga | 16260 |
| ggccaagacg | agtggatcac | ttgaggtcag | gagttccaga | tcagcctggc | caacatggta | 16320 |
| aaacccagtt | ctctactaaa | aatacaaaaa | ttagctgggc | atgggtggtga | tcgcctgtaa | 16380 |
| tcccagctac | tcaggagact | gaggcatgag | aatcgcttaa | acccaagaag | tggagggtgc | 16440 |
| agtgagctga | gatcacgcca | ctgcactcca | gcctgggcga | cagagtgaga | ctctgtctca | 16500 |
| aaaaaaaaag | aaaaaaaaaat | gttaagtga | aaagttaaga | aaccaaacaa | ggttttacaac | 16560 |
| actacatgat | ttaagcaaaa | aaaatttttt | ttgttttaga | gaaagggctc | cattctgtca | 16620 |
| tccaggcagt | gcagtgcgat | catagctctc | tgcagcctca | aactcccggg | ttcaagcagt | 16680 |
| cctcccgcct | cagcctctgg | agcagctggg | actgtaggca | cacaccacca | tgcccagcta | 16740 |
| atTTTTTgat | TTTTGTTTT | TGTAGAGACG | GGGTCTCAGT | ATGTTGCCCA | GCCTGATCTC | 16800 |
| aaactcctgg | cctcaggtga | tcctccgaag | tcagcctccc | caaagtgtctg | ggattacagg | 16860 |
| catgtgccac | catgctggcc | aattttttaa | aattttctgt | agagacaggg | tcttgctatg | 16920 |
| ttggccaggc | tggctctgaa | ctcttgacct | caagtgatcc | tgccctcaggc | tcccaaagtg | 16980 |
| atgggattac | aggcatgaac | taccacacct | ggccttaaac | ttaagcaaat | TTTTTTTTTT | 17040 |
| TTTTGGAGAC | AGTTTCACTC | TGTCGCCCAG | GCTGGAGTAA | AGTGGCGTGA | TCTCTGCTCA | 17100 |
| ctgcaacctc | cgccccccgg | gtttaagcta | ttctcctgcc | tcagcctccc | gagtagctgg | 17160 |
| gatataggcg | cctgccacca | cgctgacta | atTTTTTgtat | TTTTAGTAGA | GACGGGGTtt | 17220 |
| tgccatgttg | gccaggctgg | tctcgaactc | ctgacctcag | gcaatccgct | cccccgacc | 17280 |
| cctaccttgg | cctcccaaag | tgtaggact | acagggtgtga | gccaccatgc | ctggccaaat | 17340 |
| ttaagcaaat | gtttgaaaac | acatacccac | aggaatgtctg | cacattttac | ccagctacta | 17400 |
| tgtctagggt | cgtatctagc | acaccagcat | ggctactgtg | gagagctggg | actggatgtg | 17460 |
| agatgagagc | taaaggggaa | gtaagcaaac | caagcagggg | aaggtaagag | aagacagaag | 17520 |
| acagagagag | agggacctaa | ctctatgaga | ggagtcagac | atgtgcaatt | gaaaaagact | 17580 |
| tgctcctgtc | tctcttctgt | gaatgtttgt | gaatatccca | acgggacact | ttcacagagg | 17640 |
| agctgattga | cgtggtcaca | gccatcagcc | ttgggacacc | agaccacagt | gtgtacacta | 17700 |
| agtggcactg | atggacactt | cagcatccct | ctagctgtctg | tcccgtttcc | cctcctcggy | 17760 |
| gaccacagct | gttgccagtc | cttggtttcc | ttcaggagggy | tgtctgggta | gaccagcctg | 17820 |
| tgtgcacaca | gtccaagata | catgaacagt | gaagtgccag | gcaatccctg | caagcatggg | 17880 |
| caggtggaga | gctgaggcct | gcttgacacc | ttcctgtctc | gaagcccagt | gagcagtttc | 17940 |
| cctccctagg | gctcagtgct | atccctata | aaatggggct | tatggcagag | ctcaccacac | 18000 |
| tgggtgcatc | tggggatttg | gcgagctcat | gtgcacacca | ttgagcatgg | ggcccaacct | 18060 |
| atataaaata | ttctacgtct | gtcagctgct | gggcactgcc | actatcagcc | tcagtagtga | 18120 |
| ctgaggggaca | gggcaccagt | cagagccctg | gtgcacacag | agtgacccca | gagaagcagc | 18180 |
| cttccctctc | tgagtctctg | ttccttctgt | taggtcctga | cttcatgggt | tgttgttagc | 18240 |
| attaaggaag | tcgctggcta | atTTTTatagt | cattgaagtc | agtgggtgtc | aacctggttc | 18300 |
| ctcaaaggat | cacttccctg | aaaaaattcc | actgtccctc | ggaggcttat | gcaggccatc | 18360 |
| ccatcccttc | cctcttggtg | tgttcagctg | acagcttttt | gctcagtgag | taagtgttag | 18420 |
| gtccatttca | cagatgggct | gcaaccaagt | ttgcagtgaa | cccactaaga | ccagagctag | 18480 |
| ggccaggact | aaatgctggg | cccaatgcca | cattccctctg | tccccacacc | acatttccct | 18540 |
| catccggaga | cctgtttacc | ccaacccagg | gccccattaa | ctccctggca | gaggccctgt | 18600 |
| tacatctgct | ctgtgccacag | cctccgcccc | cccttcagga | ggcagcaggt | cccactgctg | 18660 |
| atgataaagt | tgcaggctgc | ctgagctaat | gaaggggctt | cctctaggct | gtgcacttag | 18720 |
| tcttctgctt | ccaaacccaa | tcagaggtga | ggcaccctct | ctgggcccac | ctctctcctc | 18780 |
| catttttctg | ttgggggtccc | agggaggaag | ccacttgctc | aggggccagg | aattttgcaa | 18840 |
| gcctcttgcc | ctagggagga | aggaagggag | gaggatctta | ccttgaactg | tcaagcctag | 18900 |
| agcctggtgg | ggcaggcaga | aatgggtgca | gtccatgagt | tagaaacact | agaggagaca | 18960 |
| ctttgtctgt | tggccggggc | aggcaagtta | attcccaggg | ctcctgccac | tgcatctcaa | 19020 |
| tctggaaggt | gaccaggtgg | ggcaggaccc | acgtctccca | gatgactcat | tttttctaga | 19080 |
| acaggggctt | ggctgccaaa | gaggatactt | gtttctggct | tgtggggaca | gtgggtggacc | 19140 |
| cagcatctgg | gctttatata | aagggcagct | ttgttgccct | gtaaacacac | agaccatggg | 19200 |
| tggccacttc | ttccagtaag | ttagctgggg | agttggaagt | ttaggtaaaa | ccttttgatt | 19260 |
| gacaaatgtt | ggcgaattac | catgctgtta | aatgaaacat | tgttctgcca | ccctggggct | 19320 |
| gtgggtgcct | gcgtgcaccc | tctgaaaaat | cacacaggaa | gtgggggtggg | gtctctgtga | 19380 |
| agctggtgtc | ccccagcctc | agggatgctg | cagaaatgga | atgaggacca | acagggactc | 19440 |
| agatgtccaa | ggaagctcta | cagcggagag | gacggcttgg | gaaggaggtc | caggcccagg | 19500 |
| tccctccgga | acccaatggg | tatggggcag | cctggctcct | gcctcatccc | ccttctcctg | 19560 |

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|------------|-------|
| ttgatttrtgt | cctcacagt | tatgccgccc | gcaaagcagc | cacctcaggt | gttcccagca | 19620 |
| tttatgcccc | cagcacctat | gccccacctgt | ctccccccaa | gacccccacc | ccaccagcta | 19680 |
| tgattcccat | ggggccctgcc | tacaacgggt | accctggagg | ataccctgga | gacgttgaca | 19740 |
| ggartagctc | aggtgagggc | gggggaagca | ggaacagctg | gtgggmgtgt | gctgggcatc | 19800 |
| tggacactga | ggggcagggg | ctggaaggaa | gagtgtcttg | ggagccgagg | aggggctctg | 19860 |
| ctcctgggtg | gcggccactg | acagccactc | tccccagct | ggtggccaag | gctcctatgt | 19920 |
| acccctgctt | cgggacacgg | acagcagtgt | ggcctctggg | gagaatccat | cgtcccgaag | 19980 |
| ttggatgtgc | ctgtaaggga | gaggggtggg | ccaggatcca | tcctcccaa | ccgaccacca | 20040 |
| ccccctgtc | cctagaagtc | cgcagtggct | acaggattca | ggccagccag | caggacgact | 20100 |
| ccatgcgggt | cctgtactac | atggagaagg | agctggccaa | cttcgaccct | tctcgacstg | 20160 |
| gcccccccag | tggccgtgtg | gagcggggta | agcaggagcc | ttggggctctg | agggctttta | 20220 |
| aggtgggggg | gtgaaacatg | tctccctgat | acctgccgca | gggactcttg | gtgcaaacc | 20280 |
| tggacccccg | gctcctccag | cagtacgtga | cacccccctt | ccctgcagcc | atgagtgaag | 20340 |
| tcacctccct | ccacgaggac | gactggcgat | ctcggccttc | ccggggccct | gccctcacc | 20400 |
| cgatccggga | tgaggagtgg | ggtggccact | ccccccggag | tcccagggga | tgggaccagg | 20460 |
| agcccgcag | ggagcaggca | ggcgggggct | ggcggggccag | gcggccccgg | gcccgtccg | 20520 |
| tggacgccct | ggacgacctc | accccgcga | gcaccgccga | gtcagggagc | aggtctccca | 20580 |
| cgagtaatgg | tgggaggaga | agccgggctt | acatgcccc | gcggagccgc | agccgggacg | 20640 |
| acctctatga | ccaagacgac | tcgagggact | tcccacgctc | ccgggacccc | cactacgacg | 20700 |
| acttcaggtc | tcgggagcgc | cctcctgccg | accccaggtc | ccaccaccac | cgtacccggg | 20760 |
| accctcgga | caacggctcc | aggtccgggg | acctccccta | tgatgggagg | ctactggagg | 20820 |
| aggctgtgag | gaagaagggg | tcggaggaga | ggaggagacc | ccacaaggag | gaggaggaa | 20880 |
| aggcctacta | cccgcgcgcg | ccgcccccg | actcggagac | cgactcgag | gcgtcccag | 20940 |
| agcgcaggct | caagaagggt | agggccgcgc | tccctggcgt | ccagaccgtc | cctgggcccc | 21000 |
| cagcgggtcc | ccgcggctca | taccttctt | tctttctccc | ttgcagaact | tggccctgag | 21060 |
| tcgggaaagt | ttagtcgtct | gatctgacgt | ttttctacgt | gcttttgkat | tttttttttt | 21120 |
| aatttgaagg | aacactgatg | aagccctgcc | ataccctctc | cgagtctaat | aaaacgtata | 21180 |
| atcacaagct | ctggagagaa | ccatttggtc | ggccgcgcgg | ggcgggggac | cggggctgct | 21240 |
| cccgtatgcg | tctgtaaagc | gccgcgtccc | gggggacccg | gagtccgggg | ccgggaggaa | 21300 |
| gagaccacag | ctggccccgg | ccgcgcgcgc | gccgcgcggc | ggagaacgtg | ccccgcgcag | 21360 |
| ccaccgcccc | cctgcgtgcg | cgccccggcc | ccgcccaggc | gtgcgcgatg | gccccggccc | 21420 |
| tcgcctctcg | cgcaccgcag | gctggccgcc | gggagcgcgc | gcgcgctcct | ctccccttcc | 21480 |
| agcccatccc | ccccagcccc | ccaccgacct | actttactgt | ctccaaactc | gggcagccca | 21540 |
| cctggcccc | gacgacccca | gccccgtctc | cgggtacccc | gacgttccat | ccagaccgcg | 21600 |
| gtttcaccag | ggcggcgcg | ggcgacctcg | cgccccgcgg | agccccgggc | tcgcgcgcgc | 21660 |
| ccgccccccc | ccggagacag | acagcgcgcg | cgctccccgg | ccgcctcccc | ccagcgcgcg | 21720 |
| tcgcccccg | gctcgcgcgc | ccgcgcgcgc | cgccgcgcgc | cgcgcgagc | tcaagtaaag | 21780 |
| gaggaaaaaa | aaaaggggga | aaaatagaaa | gcggcggcgg | ctgcagcagc | gatccgcgcg | 21840 |
| cggactgggc | caagccgggc | ggcggccgcg | cgagccggcg | atccagggca | ctggcgggcg | 21900 |
| ccagccaggg | cgggcccgtg | tcaaaaaaaa | aagtcgcggc | ggcggcggtt | gctcagggaa | 21960 |
| ggaggccctga | gggcccgcgtg | cagcgggcgg | gcagctgggt | gggctggggg | cgccgcgcgc | 22020 |
| gcgtcccga | gcctcggggc | gcccggagcc | ggcgggcggg | cgagggcgga | ggcggcgggc | 22080 |
| gctgcagcgg | ctgcaggagc | ggcggcggtt | gcngcggcgg | cngcggcatc | tcctcctcac | 22140 |
| atgacccac | tgtttgtccc | cgtgatcagc | gcgagcggct | cccgtatctc | ctccgtcccc | 22200 |
| tcctgcgcgc | cggcgtgagc | gcccgggctc | ggggcccccc | cgccgcgcgc | ccccctcccc | 22260 |
| tcctccctc | ccctccctcc | ccctccctcc | cgggccccgc | gccccccccg | ccccgccccc | 22320 |
| ccccatggac | atgctggacc | cgggtctgga | tcccgctgcc | tcggccaccg | ctgctgcgcg | 22380 |
| cgccaggtaa | gatccccggc | ccggccgtgc | ccccgcgcgc | cggccccggc | cccgcccccg | 22440 |
| cggcctgcag | gcccggggccg | ccatgatccc | gagcgggcgc | gggccccgct | caaaatggag | 22500 |
| gcccgcggcg | cggggggggac | ctggcgccctc | ccgcccccg | ccccggcct | cggcggcgcc | 22560 |
| ccgggcctca | ggcgcggccg | ggtgggactg | ggccctcgca | gctgggcgcg | ggggcggggg | 22620 |
| cgcgggcgcg | ggcgcgcgtg | accctgctcc | ctcctgtgcc | cctggcagcc | acgacaaggg | 22680 |
| acccgaggcg | gaggaggcg | tcgagctgca | ggaagggtag | tgcttgccgg | gcccgcgcgc | 22740 |
| ccgggggagg | gctggggggc | ctcggcgcg | ccctgaccgt | gccccgaccc | tcctcgggcc | 22800 |
| caggcgggga | cggcccaggga | gcggaggagc | agacagcggt | ggccatcacc | agcgtccagc | 22860 |
| aggcggcggt | cggcgaccac | aacatccagt | accagttccg | cacagagaca | aatggaggag | 22920 |
| aggtgagcgg | cgggcccgcga | gagcgaacgg | gcgggcgggc | gggcgcgcgc | ggaagggtcg | 22980 |
| gacctggccc | cagcgcgggc | ctcgcgcgtc | tgccgcccc | tgcaggtgac | ataccgcgta | 23040 |
| gtccagggtga | ctgatgggtca | gctggacggc | cagggcgaca | cagctggcgc | cgtcagcgtc | 23100 |

gtgtccaccg ctgccttcgc ggggggggcag caggctgtga cccagggtggg tgtggacggg 23160
gcagcccagc gcccgggccc cgccgct 23187

<210> 2
<211> 2158
<212> DNA
<213> Homo sapiens

<220>
<221> allele
<222> 595
<223> 9-3-324 : polymorphic base C or T

<220>
<221> allele
<222> 940
<223> 9-6-187 : polymorphic base C or T

<220>
<221> allele
<222> 1191
<223> 9-7-325 : polymorphic base A or G

<220>
<221> allele
<222> 1362
<223> 9-9-246 : polymorphic base G or C

<220>
<221> allele
<222> 1658
<223> LSRX9f13-BM : polymorphic base deletion of AGG

<220>
<221> allele
<222> 2079
<223> LSRX9f14-BM : polymorphic base T or G

<400> 2
tggagtgtgg ctgcggaggac cgcgggcgggt caagcacctt tctcccccat atctgaaagc 60
atgccctttg tccacgtcgt ttacgctcat taaaacttcc aga atg caa cag gac 115
Met Gln Gln Asp
1
gga ctt gga gta ggg aca agg aac gga agt ggg aag ggg agg agc gtg 163
Gly Leu Gly Val Gly Thr Arg Asn Gly Ser Gly Lys Gly Arg Ser Val
5 10 15 20
cac ccc tcc tgg cct tgg tgc gcg ccg cgc ccc cta agg tac ttt gga 211
His Pro Ser Trp Pro Trp Cys Ala Pro Arg Pro Leu Arg Tyr Phe Gly
25 30 35
agg gac gcg cgg gcc aga cgc gcc cag acg gcc gcg atg gcg ctg ttg 259
Arg Asp Ala Arg Ala Arg Arg Ala Gln Thr Ala Ala Met Ala Leu Leu
40 45 50
gcc ggc ggg ctc tcc aga ggg ctg ggc tcc cac ccg gcc gcc gca ggc 307
Ala Gly Gly Leu Ser Arg Gly Leu Gly Ser His Pro Ala Ala Ala Gly
55 60 65
cgg gac gcg gtc gtc ttc gtg tgg ctt ctg ctt agc acc tgg tgc aca 355
Arg Asp Ala Val Val Phe Val Trp Leu Leu Leu Ser Thr Trp Cys Thr
70 75 80
gct cct gcc agg gcc atc cag gtg acc gtg tcc aac ccc tac cac gtg 403

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|--|
| Ala | Pro | Ala | Arg | Ala | Ile | Gln | Val | Thr | Val | Ser | Asn | Pro | Tyr | His | Val | | |
| 85 | | | | | 90 | | | | | 95 | | | | | 100 | | |
| gtg | atc | ctc | ttc | cag | cct | gtg | acc | ctg | ccc | tgt | acc | tac | cag | atg | acc | 451 | |
| Val | Ile | Leu | Phe | Gln | Pro | Val | Thr | Leu | Pro | Cys | Thr | Tyr | Gln | Met | Thr | | |
| | | | | 105 | | | | | 110 | | | | | 115 | | | |
| tcg | acc | ccc | acg | caa | ccc | atc | gtc | atc | tgg | aag | tac | aag | tct | ttc | tgc | 499 | |
| Ser | Thr | Pro | Thr | Gln | Pro | Ile | Val | Ile | Trp | Lys | Tyr | Lys | Ser | Phe | Cys | | |
| | | | 120 | | | | | 125 | | | | | 130 | | | | |
| cgg | gac | cgc | atc | gcc | gat | gcc | ttc | tcc | ccg | gcc | agc | gtc | gac | aac | cag | 547 | |
| Arg | Asp | Arg | Ile | Ala | Asp | Ala | Phe | Ser | Pro | Ala | Ser | Val | Asp | Asn | Gln | | |
| | | | 135 | | | | 140 | | | | | 145 | | | | | |
| ctc | aat | gcc | cag | ctg | gca | gcc | ggg | aac | cca | ggc | tac | aac | ccc | tac | gty | 595 | |
| Leu | Asn | Ala | Gln | Leu | Ala | Ala | Gly | Asn | Pro | Gly | Tyr | Asn | Pro | Tyr | Val | | |
| | 150 | | | | | 155 | | | | 160 | | | | | | | |
| gag | tgc | cag | gac | agc | gtg | cgc | acc | gtc | agg | gtc | gtg | gcc | acc | aag | cag | 643 | |
| Glu | Cys | Gln | Asp | Ser | Val | Arg | Thr | Val | Arg | Val | Val | Ala | Thr | Lys | Gln | | |
| | 165 | | | | 170 | | | | 175 | | | | | | 180 | | |
| ggc | aac | gct | gtg | acc | ctg | gga | gat | tac | tac | cag | ggc | cgg | agg | att | acc | 691 | |
| Gly | Asn | Ala | Val | Thr | Leu | Gly | Asp | Tyr | Tyr | Gln | Gly | Arg | Arg | Ile | Thr | | |
| | | | 185 | | | | | 190 | | | | | | 195 | | | |
| atc | acc | gga | aat | gct | gac | ctg | acc | ttt | gac | cag | acg | gcg | tgg | ggg | gac | 739 | |
| Ile | Thr | Gly | Asn | Ala | Asp | Leu | Thr | Phe | Asp | Gln | Thr | Ala | Trp | Gly | Asp | | |
| | | | 200 | | | | | 205 | | | | | 210 | | | | |
| agt | ggt | gtg | tat | tac | tgc | tcc | gtg | gtc | tca | gcc | cag | gac | ctc | cag | ggg | 787 | |
| Ser | Gly | Val | Tyr | Tyr | Cys | Ser | Val | Val | Ser | Ala | Gln | Asp | Leu | Gln | Gly | | |
| | | 215 | | | | | 220 | | | | | 225 | | | | | |
| aac | aat | gag | gcc | tac | gca | gag | ctc | atc | gtc | ctt | ggg | agg | acc | tca | ggg | 835 | |
| Asn | Asn | Glu | Ala | Tyr | Ala | Glu | Leu | Ile | Val | Leu | Gly | Arg | Thr | Ser | Gly | | |
| | | 230 | | | | 235 | | | | 240 | | | | | | | |
| gtg | gct | gag | ctc | tta | cct | ggt | ttt | cag | gcg | ggg | ccc | ata | gaa | gac | tgg | 883 | |
| Val | Ala | Glu | Leu | Leu | Pro | Gly | Phe | Gln | Ala | Gly | Pro | Ile | Glu | Asp | Trp | | |
| | 245 | | | | 250 | | | | 255 | | | | | | 260 | | |
| ctc | ttc | gtg | gtt | gtg | gta | tgc | ctg | gct | gcc | ttc | ctc | atc | ttc | ctc | ctc | 931 | |
| Leu | Phe | Val | Val | Val | Val | Cys | Leu | Ala | Ala | Phe | Leu | Ile | Phe | Leu | Leu | | |
| | | | 265 | | | | | 270 | | | | | | 275 | | | |
| ctg | ggc | aty | tgc | tgg | tgc | cag | tgc | tgc | ccg | cac | act | tgc | tgc | tgc | tac | 979 | |
| Leu | Gly | Ile | Cys | Trp | Cys | Gln | Cys | Cys | Pro | His | Thr | Cys | Cys | Cys | Tyr | | |
| | | 280 | | | | | 285 | | | | | | 290 | | | | |
| gtc | agg | tgc | ccc | tgc | tgc | cca | gac | aag | tgc | tgc | tgc | ccc | gag | gcc | ctg | 1027 | |
| Val | Arg | Cys | Pro | Cys | Cys | Pro | Asp | Lys | Cys | Cys | Cys | Pro | Glu | Ala | Leu | | |
| | | 295 | | | | 300 | | | | | | 305 | | | | | |
| tat | gcc | gcc | ggc | aaa | gca | gcc | acc | tca | ggt | gtt | ccc | agc | att | tat | gcc | 1075 | |
| Tyr | Ala | Ala | Gly | Lys | Ala | Ala | Thr | Ser | Gly | Val | Pro | Ser | Ile | Tyr | Ala | | |
| | 310 | | | | 315 | | | | | 320 | | | | | | | |
| ccc | agc | acc | tat | gcc | cac | ctg | tct | ccc | gcc | aag | acc | cca | ccc | cca | cca | 1123 | |
| Pro | Ser | Thr | Tyr | Ala | His | Leu | Ser | Pro | Ala | Lys | Thr | Pro | Pro | Pro | Pro | | |
| | 325 | | | 330 | | | | | 335 | | | | | | 340 | | |
| gct | atg | att | ccc | atg | ggc | cct | gcc | tac | aac | ggg | tac | cct | gga | gga | tac | 1171 | |
| Ala | Met | Ile | Pro | Met | Gly | Pro | Ala | Tyr | Asn | Gly | Tyr | Pro | Gly | Gly | Tyr | | |
| | | | 345 | | | | | 350 | | | | | 355 | | | | |
| cct | gga | gac | gtt | gac | agg | art | agc | tca | gct | ggt | ggc | caa | ggc | tcc | tat | 1219 | |
| Pro | Gly | Asp | Val | Asp | Arg | Xaa | Ser | Ser | Ala | Gly | Gly | Gln | Gly | Ser | Tyr | | |
| | | 360 | | | | | 365 | | | | | 370 | | | | | |
| gta | ccc | ctg | ctt | cgg | gac | acg | gac | agc | agt | gtg | gcc | tct | gaa | gtc | cgc | 1267 | |
| Val | Pro | Leu | Leu | Arg | Asp | Thr | Asp | Ser | Ser | Val | Ala | Ser | Glu | Val | Arg | | |
| | | 375 | | | | 380 | | | | | 385 | | | | | | |
| agt | ggc | tac | agg | att | cag | gcc | agc | cag | cag | gac | gac | tcc | atg | cgg | gtc | 1315 | |
| Ser | Gly | Tyr | Arg | Ile | Gln | Ala | Ser | Gln | Gln | Asp | Asp | Ser | Met | Arg | Val | | |
| | 390 | | | | 395 | | | | | 400 | | | | | | | |

| | |
|---|------|
| ctg tac tac atg gag aag gag ctg gcc aac ttc gac cct tct cga cst | 1363 |
| Leu Tyr Tyr Met Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser Arg Xaa | |
| 405 410 415 420 | |
| ggc ccc ccc agt ggc cgt gtg gag cgg gcc atg agt gaa gtc acc tcc | 1411 |
| Gly Pro Pro Ser Gly Arg Val Glu Arg Ala Met Ser Glu Val Thr Ser | |
| 425 430 435 | |
| ctc cac gag gac gac tgg cga tct cgg cct tcc cgg ggc cct gcc ctc | 1459 |
| Leu His Glu Asp Asp Trp Arg Ser Arg Pro Ser Arg Gly Pro Ala Leu | |
| 440 445 450 | |
| acc ccg atc cgg gat gag gag tgg ggt ggc cac tcc ccc cgg agt ccc | 1507 |
| Thr Pro Ile Arg Asp Glu Glu Trp Gly Gly His Ser Pro Arg Ser Pro | |
| 455 460 465 | |
| agg gga tgg gac cag gag ccc gcc agg gag cag gca ggc ggg ggc tgg | 1555 |
| Arg Gly Trp Asp Gln Glu Pro Ala Arg Glu Gln Ala Gly Gly Gly Trp | |
| 470 475 480 | |
| cgg gcc agg cgg ccc cgg gcc cgc tcc gtg gac gcc ctg gac gac ctc | 1603 |
| Arg Ala Arg Arg Pro Arg Ala Arg Ser Val Asp Ala Leu Asp Asp Leu | |
| 485 490 495 500 | |
| acc ccg ccg agc acc gcc gag tca ggg agc agg tct ccc acg agt aat | 1651 |
| Thr Pro Pro Ser Thr Ala Glu Ser Gly Ser Arg Ser Pro Thr Ser Asn | |
| 505 510 515 | |
| ggt ggg aga agc cgg gcc tac atg ccc ccg cgg agc cgc agc cgg gac | 1699 |
| Gly Gly Arg Ser Arg Ala Tyr Met Pro Pro Arg Ser Arg Ser Arg Asp | |
| 520 525 530 | |
| gac ctc tat gac caa gac gac tcg agg gac ttc cca cgc tcc cgg gac | 1747 |
| Asp Leu Tyr Asp Gln Asp Asp Ser Arg Asp Phe Pro Arg Ser Arg Asp | |
| 535 540 545 | |
| ccc cac tac gac gac ttc agg tct cgg gag cgc cct cct gcc gac ccc | 1795 |
| Pro His Tyr Asp Asp Phe Arg Ser Arg Glu Arg Pro Pro Ala Asp Pro | |
| 550 555 560 | |
| agg tcc cac cac cac cgt acc cgg gac cct cgg gac aac ggc tcc agg | 1843 |
| Arg Ser His His His Arg Thr Arg Asp Pro Arg Asp Asn Gly Ser Arg | |
| 565 570 575 580 | |
| tcc ggg gac ctc ccc tat gat ggg cgg cta ctg gag gag gct gtg agg | 1891 |
| Ser Gly Asp Leu Pro Tyr Asp Gly Arg Leu Leu Glu Glu Ala Val Arg | |
| 585 590 595 | |
| aag aag ggg tcg gag gag agg agg aga ccc cac aag gag gag gag gaa | 1939 |
| Lys Lys Gly Ser Glu Glu Arg Arg Arg Pro His Lys Glu Glu Glu Glu | |
| 600 605 610 | |
| gag gcc tac tac ccg ccc gcg ccg ccc ccg tac tcg gag acc gac tcg | 1987 |
| Glu Ala Tyr Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp Ser | |
| 615 620 625 | |
| cag gcg tcc cga gag cgc agg ctc aag aag aac ttg gcc ctg agt cgg | 2035 |
| Gln Ala Ser Arg Glu Arg Arg Leu Lys Lys Asn Leu Ala Leu Ser Arg | |
| 630 635 640 | |
| gaa agt tta gtc gtc tga tctgacgttt tctacgtagc ttttgkattt | 2083 |
| Glu Ser Leu Val Val * | |
| 645 650 | |
| ttttttttaa tttgaaggaa cactgatgaa gccctgccat acccctcccg agtctaataa | 2143 |
| aacgtataat caca | 2158 |

<210> 3
 <211> 649
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> 363

<223> 9-7-325 : polymorphic amino acid Ser or Asn

<220>

<221> VARIANT

<222> 420

<223> 9-9-246 : polymorphic amino acid Pro or Arg

<220>

<221> VARIANT

<222> 519

<223> LSRX9f13-BM : polymorphic amino acid deletion of Arg

<400> 3

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gln | Gln | Asp | Gly | Leu | Gly | Val | Gly | Thr | Arg | Asn | Gly | Ser | Gly | Lys |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Gly | Arg | Ser | Val | His | Pro | Ser | Trp | Pro | Trp | Cys | Ala | Pro | Arg | Pro | Leu |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Arg | Tyr | Phe | Gly | Arg | Asp | Ala | Arg | Ala | Arg | Arg | Ala | Gln | Thr | Ala | Ala |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| Met | Ala | Leu | Leu | Ala | Gly | Gly | Leu | Ser | Arg | Gly | Leu | Gly | Ser | His | Pro |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| Ala | Ala | Ala | Gly | Arg | Asp | Ala | Val | Val | Phe | Val | Trp | Leu | Leu | Leu | Ser |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| Thr | Trp | Cys | Thr | Ala | Pro | Ala | Arg | Ala | Ile | Gln | Val | Thr | Val | Ser | Asn |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| Pro | Tyr | His | Val | Val | Ile | Leu | Phe | Gln | Pro | Val | Thr | Leu | Pro | Cys | Thr |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Tyr | Gln | Met | Thr | Ser | Thr | Pro | Thr | Gln | Pro | Ile | Val | Ile | Trp | Lys | Tyr |
| | | 115 | | | | | 120 | | | | | 125 | | | |
| Lys | Ser | Phe | Cys | Arg | Asp | Arg | Ile | Ala | Asp | Ala | Phe | Ser | Pro | Ala | Ser |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| Val | Asp | Asn | Gln | Leu | Asn | Ala | Gln | Leu | Ala | Ala | Gly | Asn | Pro | Gly | Tyr |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Asn | Pro | Tyr | Val | Glu | Cys | Gln | Asp | Ser | Val | Arg | Thr | Val | Arg | Val | Val |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Ala | Thr | Lys | Gln | Gly | Asn | Ala | Val | Thr | Leu | Gly | Asp | Tyr | Tyr | Gln | Gly |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Arg | Arg | Ile | Thr | Ile | Thr | Gly | Asn | Ala | Asp | Leu | Thr | Phe | Asp | Gln | Thr |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Ala | Trp | Gly | Asp | Ser | Gly | Val | Tyr | Tyr | Cys | Ser | Val | Val | Ser | Ala | Gln |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Asp | Leu | Gln | Gly | Asn | Asn | Glu | Ala | Tyr | Ala | Glu | Leu | Ile | Val | Leu | Gly |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Arg | Thr | Ser | Gly | Val | Ala | Glu | Leu | Leu | Pro | Gly | Phe | Gln | Ala | Gly | Pro |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Ile | Glu | Asp | Trp | Leu | Phe | Val | Val | Val | Val | Cys | Leu | Ala | Ala | Phe | Leu |
| | | 260 | | | | | | 265 | | | | | 270 | | |
| Ile | Phe | Leu | Leu | Leu | Gly | Ile | Cys | Trp | Cys | Gln | Cys | Cys | Pro | His | Thr |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Cys | Cys | Cys | Tyr | Val | Arg | Cys | Pro | Cys | Cys | Pro | Asp | Lys | Cys | Cys | Cys |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Pro | Glu | Ala | Leu | Tyr | Ala | Ala | Gly | Lys | Ala | Ala | Thr | Ser | Gly | Val | Pro |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Ser | Ile | Tyr | Ala | Pro | Ser | Thr | Tyr | Ala | His | Leu | Ser | Pro | Ala | Lys | Thr |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Pro | Pro | Pro | Pro | Ala | Met | Ile | Pro | Met | Gly | Pro | Ala | Tyr | Asn | Gly | Tyr |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| Pro | Gly | Gly | Tyr | Pro | Gly | Asp | Val | Asp | Arg | Ser | Ser | Ser | Ala | Gly | Gly |
| | | 355 | | | | | 360 | | | | | | 365 | | |

Gln Gly Ser Tyr Val Pro Leu Leu Arg Asp Thr Asp Ser Ser Val Ala
 370 375 380
 Ser Glu Val Arg Ser Gly Tyr Arg Ile Gln Ala Ser Gln Gln Asp Asp
 385 390 395 400
 Ser Met Arg Val Leu Tyr Tyr Met Glu Lys Glu Leu Ala Asn Phe Asp
 405 410 415
 Pro Ser Arg Pro Gly Pro Pro Ser Gly Arg Val Glu Arg Ala Met Ser
 420 425 430
 Glu Val Thr Ser Leu His Glu Asp Asp Trp Arg Ser Arg Pro Ser Arg
 435 440 445
 Gly Pro Ala Leu Thr Pro Ile Arg Asp Glu Glu Trp Gly Gly His Ser
 450 455 460
 Pro Arg Ser Pro Arg Gly Trp Asp Gln Glu Pro Ala Arg Glu Gln Ala
 465 470 475 480
 Gly Gly Gly Trp Arg Ala Arg Arg Pro Arg Ala Arg Ser Val Asp Ala
 485 490 495
 Leu Asp Asp Leu Thr Pro Pro Ser Thr Ala Glu Ser Gly Ser Arg Ser
 500 505 510
 Pro Thr Ser Asn Gly Gly Arg Ser Arg Ala Tyr Met Pro Pro Arg Ser
 515 520 525
 Arg Ser Arg Asp Asp Leu Tyr Asp Gln Asp Asp Ser Arg Asp Phe Pro
 530 535 540
 Arg Ser Arg Asp Pro His Tyr Asp Asp Phe Arg Ser Arg Glu Arg Pro
 545 550 555 560
 Pro Ala Asp Pro Arg Ser His His His Arg Thr Arg Asp Pro Arg Asp
 565 570 575
 Asn Gly Ser Arg Ser Gly Asp Leu Pro Tyr Asp Gly Arg Leu Leu Glu
 580 585 590
 Glu Ala Val Arg Lys Lys Gly Ser Glu Glu Arg Arg Arg Pro His Lys
 595 600 605
 Glu Glu Glu Glu Glu Ala Tyr Pro Pro Ala Pro Pro Pro Tyr Ser
 610 615 620
 Glu Thr Asp Ser Gln Ala Ser Arg Glu Arg Arg Leu Lys Lys Asn Leu
 625 630 635 640
 Ala Leu Ser Arg Glu Ser Leu Val Val
 645

<210> 4
 <211> 2101
 <212> DNA
 <213> Homo sapiens

<220>
 <221> allele
 <222> 595
 <223> 9-3-324 : polymorphic base C or T

<220>
 <221> allele
 <222> 883
 <223> 9-6-187 : polymorphic base C or T

<220>
 <221> allele
 <222> 1134
 <223> 9-7-325 : polymorphic base A or G

<220>
 <221> allele

<222> 1305
 <223> 9-9-246 : polymorphic base G or C

<220>
 <221> allele
 <222> 1601
 <223> LSRX9f13-BM : polymorphic base deletion of AGG

<220>
 <221> allele
 <222> 2022
 <223> LSRX9f14-BM : polymorphic base T or G

<400> 4
 tggagtgtgg ctcggaggac cgcggcgggt caagcacctt tctcccccat atctgaaagc 60
 atgccctttg tccacgtcgt ttacgctcat taaaacttcc aga atg caa cag gac 115
 Met Gln Gln Asp
 1
 gga ctt gga gta ggg aca agg aac gga agt ggg aag ggg agg agc gtg 163
 Gly Leu Gly Val Gly Thr Arg Asn Gly Ser Gly Lys Gly Arg Ser Val
 5 10 15 20
 cac ccc tcc tgg cct tgg tgc gcg ccg cgc ccc cta agg tac ttt gga 211
 His Pro Ser Trp Pro Trp Cys Ala Pro Arg Pro Leu Arg Tyr Phe Gly
 25 30 35
 agg gac gcg cgg gcc aga cgc gcc cag acg gcc gcg atg gcg ctg ttg 259
 Arg Asp Ala Arg Ala Arg Arg Ala Gln Thr Ala Ala Met Ala Leu Leu
 40 45 50
 gcc ggc ggg ctc tcc aga ggg ctg ggc tcc cac ccg gcc gcc gca ggc 307
 Ala Gly Gly Leu Ser Arg Gly Leu Gly Ser His Pro Ala Ala Ala Gly
 55 60 65
 cgg gac gcg gtc gtc ttc gtg tgg ctt ctg ctt agc acc tgg tgc aca 355
 Arg Asp Ala Val Val Phe Val Trp Leu Leu Leu Ser Thr Trp Cys Thr
 70 75 80
 gct cct gcc agg gcc atc cag gtg acc gtg tcc aac ccc tac cac gtg 403
 Ala Pro Ala Arg Ala Ile Gln Val Thr Val Ser Asn Pro Tyr His Val
 85 90 95 100
 gtg atc ctc ttc cag cct gtg acc ctg ccc tgt acc tac cag atg acc 451
 Val Ile Leu Phe Gln Pro Val Thr Leu Pro Cys Thr Tyr Gln Met Thr
 105 110 115
 tcg acc ccc acg caa ccc atc gtc atc tgg aag tac aag tct ttc tgc 499
 Ser Thr Pro Thr Gln Pro Ile Val Ile Trp Lys Tyr Lys Ser Phe Cys
 120 125 130
 cgg gac cgc atc gcc gat gcc ttc tcc ccg gcc agc gtc gac aac cag 547
 Arg Asp Arg Ile Ala Asp Ala Phe Ser Pro Ala Ser Val Asp Asn Gln
 135 140 145
 ctc aat gcc cag ctg gca gcc ggg aac cca ggc tac aac ccc tac gty 595
 Leu Asn Ala Gln Leu Ala Ala Gly Asn Pro Gly Tyr Asn Pro Tyr Val
 150 155 160
 gag tgc cag gac agc gtg cgc acc gtc agg gtc gtg gcc acc aag cag 643
 Glu Cys Gln Asp Ser Val Arg Thr Val Arg Val Val Ala Thr Lys Gln
 165 170 175 180
 ggc aac gct gtg acc ctg gga gat tac tac cag ggc cgg agg att acc 691
 Gly Asn Ala Val Thr Leu Gly Asp Tyr Tyr Gln Gly Arg Arg Ile Thr
 185 190 195
 atc acc gga aat gct gac ctg acc ttt gac cag acg gcg tgg ggg gac 739
 Ile Thr Gly Asn Ala Asp Leu Thr Phe Asp Gln Thr Ala Trp Gly Asp
 200 205 210
 agt ggt gtg tat tac tgc tcc gtg gtc tca gcc cag gac ctc cag ggg 787
 Ser Gly Val Tyr Tyr Cys Ser Val Val Ser Ala Gln Asp Leu Gln Gly

| | | | |
|---|-----|-----|------|
| 215 | 220 | 225 | |
| aac aat gag gcc tac gca gag ctc atc gtc ctt gac tgg ctc ttc gtg | | | 835 |
| Asn Asn Glu Ala Tyr Ala Glu Leu Ile Val Leu Asp Trp Leu Phe Val | | | |
| 230 | 235 | 240 | |
| gtt gtg gta tgc ctg gct gcc ttc ctc atc ttc ctc ctc ctg ggc aty | | | 883 |
| Val Val Val Cys Leu Ala Ala Phe Leu Ile Phe Leu Leu Leu Gly Ile | | | |
| 245 | 250 | 255 | 260 |
| tgc tgg tgc cag tgc tgc ccg cac act tgc tgc tgc tac gtc agg tgc | | | 931 |
| Cys Trp Cys Gln Cys Cys Pro His Thr Cys Cys Cys Tyr Val Arg Cys | | | |
| 265 | 270 | 275 | |
| ccc tgc tgc cca gac aag tgc tgc tgc ccc gag gcc ctg tat gcc gcc | | | 979 |
| Pro Cys Cys Pro Asp Lys Cys Cys Cys Pro Glu Ala Leu Tyr Ala Ala | | | |
| 280 | 285 | 290 | |
| ggc aaa gca gcc acc tca ggt gtt ccc agc att tat gcc ccc agc acc | | | 1027 |
| Gly Lys Ala Ala Thr Ser Gly Val Pro Ser Ile Tyr Ala Pro Ser Thr | | | |
| 295 | 300 | 305 | |
| tat gcc cac ctg tct ccc gcc aag acc cca ccc cca cca gct atg att | | | 1075 |
| Tyr Ala His Leu Ser Pro Ala Lys Thr Pro Pro Pro Pro Ala Met Ile | | | |
| 310 | 315 | 320 | |
| ccc atg ggc cct gcc tac aac ggg tac cct gga gga tac cct gga gac | | | 1123 |
| Pro Met Gly Pro Ala Tyr Asn Gly Tyr Pro Gly Gly Tyr Pro Gly Asp | | | |
| 325 | 330 | 335 | 340 |
| gtt gac agg art agc tca gct ggt ggc caa ggc tcc tat gta ccc ctg | | | 1171 |
| Val Asp Arg Xaa Ser Ser Ala Gly Gly Gln Gly Ser Tyr Val Pro Leu | | | |
| 345 | 350 | 355 | |
| ctt cgg gac acg gac agc agt gtg gcc tct gaa gtc cgc agt ggc tac | | | 1219 |
| Leu Arg Asp Thr Asp Ser Ser Val Ala Ser Glu Val Arg Ser Gly Tyr | | | |
| 360 | 365 | 370 | |
| agg att cag gcc agc cag cag gac gac tcc atg cgg gtc ctg tac tac | | | 1267 |
| Arg Ile Gln Ala Ser Gln Gln Asp Asp Ser Met Arg Val Leu Tyr Tyr | | | |
| 375 | 380 | 385 | |
| atg gag aag gag ctg gcc aac ttc gac cct tct cga cst ggc ccc ccc | | | 1315 |
| Met Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser Arg Xaa Gly Pro Pro | | | |
| 390 | 395 | 400 | |
| agt ggc cgt gtg gag cgg gcc atg agt gaa gtc acc tcc ctc cac gag | | | 1363 |
| Ser Gly Arg Val Glu Arg Ala Met Ser Glu Val Thr Ser Leu His Glu | | | |
| 405 | 410 | 415 | 420 |
| gac gac tgg cga tct cgg cct tcc cgg ggc cct gcc ctc acc ccg atc | | | 1411 |
| Asp Asp Trp Arg Ser Arg Pro Ser Arg Gly Pro Ala Leu Thr Pro Ile | | | |
| 425 | 430 | 435 | |
| cgg gat gag gag tgg ggt ggc cac tcc ccc cgg agt ccc agg gga tgg | | | 1459 |
| Arg Asp Glu Glu Trp Gly Gly His Ser Pro Arg Ser Pro Arg Gly Trp | | | |
| 440 | 445 | 450 | |
| gac cag gag ccc gcc agg gag cag gca ggc ggg ggc tgg cgg gcc agg | | | 1507 |
| Asp Gln Glu Pro Ala Arg Glu Gln Ala Gly Gly Gly Trp Arg Ala Arg | | | |
| 455 | 460 | 465 | |
| cgg ccc cgg gcc cgc tcc gtg gac gcc ctg gac gac ctc acc ccg ccg | | | 1555 |
| Arg Pro Arg Ala Arg Ser Val Asp Ala Leu Asp Asp Leu Thr Pro Pro | | | |
| 470 | 475 | 480 | |
| agc acc gcc gag tca ggg agc agg tct ccc acg agt aat ggt ggg aga | | | 1603 |
| Ser Thr Ala Glu Ser Gly Ser Arg Ser Pro Thr Ser Asn Gly Gly Arg | | | |
| 485 | 490 | 495 | 500 |
| agc cgg gcc tac atg ccc ccg cgg agc cgc agc cgg gac gac ctc tat | | | 1651 |
| Ser Arg Ala Tyr Met Pro Pro Arg Ser Arg Ser Arg Asp Asp Leu Tyr | | | |
| 505 | 510 | 515 | |
| gac caa gac gac tcc agg gac ttc cca cgc tcc cgg gac ccc cac tac | | | 1699 |
| Asp Gln Asp Asp Ser Arg Asp Phe Pro Arg Ser Arg Asp Pro His Tyr | | | |
| 520 | 525 | 530 | |
| gac gac ttc agg tct cgg gag cgc cct cct gcc gac ccc agg tcc cac | | | 1747 |

| | |
|---|------|
| Asp Asp Phe Arg Ser Arg Glu Arg Pro Pro Ala Asp Pro Arg Ser His | |
| 535 540 545 | |
| cac cac cgt acc cgg gac cct cgg gac aac ggc tcc agg tcc ggg gac | 1795 |
| His His Arg Thr Arg Asp Pro Arg Asp Asn Gly Ser Arg Ser Gly Asp | |
| 550 555 560 | |
| ctc ccc tat gat ggg cgg cta ctg gag gag gct gtg agg aag aag ggg | 1843 |
| Leu Pro Tyr Asp Gly Arg Leu Leu Glu Glu Ala Val Arg Lys Lys Gly | |
| 565 570 575 580 | |
| tcg gag gag agg agg aga ccc cac aag gag gag gag gaa gag gcc tac | 1891 |
| Ser Glu Glu Arg Arg Arg Pro His Lys Glu Glu Glu Glu Glu Ala Tyr | |
| 585 590 595 | |
| tac ccg ccc gcg ccg ccc ccg tac tcg gag acc gac tcg cag gcg tcc | 1939 |
| Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp Ser Gln Ala Ser | |
| 600 605 610 | |
| cga gag cgc agg ctc aag aag aac ttg gcc ctg agt cgg gaa agt tta | 1987 |
| Arg Glu Arg Arg Leu Lys Lys Asn Leu Ala Leu Ser Arg Glu Ser Leu | |
| 615 620 625 | |
| gtc gtc tga tctgacgttt tctacgtagc ttttgkattt ttttttttaa | 2036 |
| Val Val * | |
| 630 | |
| tttgaaggaa cactgatgaa gccctgccat acccctcccg agtctaataa aacgtataat | 2096 |
| cacaa | 2101 |

<210> 5
 <211> 630
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> 344
 <223> 9-7-325 : polymorphic amino acid Ser or Asn

<220>
 <221> VARIANT
 <222> 401
 <223> 9-9-246 : polymorphic amino acid Pro or Arg

<220>
 <221> VARIANT
 <222> 500
 <223> LSRX9f13-BM : polymorphic amino acid deletion of Arg

| | |
|---|--|
| <400> 5 | |
| Met Gln Gln Asp Gly Leu Gly Val Gly Thr Arg Asn Gly Ser Gly Lys | |
| 1 5 10 15 | |
| Gly Arg Ser Val His Pro Ser Trp Pro Trp Cys Ala Pro Arg Pro Leu | |
| 20 25 30 | |
| Arg Tyr Phe Gly Arg Asp Ala Arg Ala Arg Arg Ala Gln Thr Ala Ala | |
| 35 40 45 | |
| Met Ala Leu Leu Ala Gly Gly Leu Ser Arg Gly Leu Gly Ser His Pro | |
| 50 55 60 | |
| Ala Ala Ala Gly Arg Asp Ala Val Val Phe Val Trp Leu Leu Leu Ser | |
| 65 70 75 80 | |
| Thr Trp Cys Thr Ala Pro Ala Arg Ala Ile Gln Val Thr Val Ser Asn | |
| 85 90 95 | |
| Pro Tyr His Val Val Ile Leu Phe Gln Pro Val Thr Leu Pro Cys Thr | |
| 100 105 110 | |
| Tyr Gln Met Thr Ser Thr Pro Thr Gln Pro Ile Val Ile Trp Lys Tyr | |

| | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| | | 115 | | | | | 120 | | | | | 125 | | | | |
| Lys | Ser | Phe | Cys | Arg | Asp | Arg | Ile | Ala | Asp | Ala | Phe | Ser | Pro | Ala | Ser | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Val | Asp | Asn | Gln | Leu | Asn | Ala | Gln | Leu | Ala | Ala | Gly | Asn | Pro | Gly | Tyr | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | |
| Asn | Pro | Tyr | Val | Glu | Cys | Gln | Asp | Ser | Val | Arg | Thr | Val | Arg | Val | Val | |
| | | | | 165 | | | | | 170 | | | | | 175 | | |
| Ala | Thr | Lys | Gln | Gly | Asn | Ala | Val | Thr | Leu | Gly | Asp | Tyr | Tyr | Gln | Gly | |
| | | | 180 | | | | | 185 | | | | | 190 | | | |
| Arg | Arg | Ile | Thr | Ile | Thr | Gly | Asn | Ala | Asp | Leu | Thr | Phe | Asp | Gln | Thr | |
| | | 195 | | | | | 200 | | | | | 205 | | | | |
| Ala | Trp | Gly | Asp | Ser | Gly | Val | Tyr | Tyr | Cys | Ser | Val | Val | Ser | Ala | Gln | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Asp | Leu | Gln | Gly | Asn | Asn | Glu | Ala | Tyr | Ala | Glu | Leu | Ile | Val | Leu | Asp | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | |
| Trp | Leu | Phe | Val | Val | Val | Val | Cys | Leu | Ala | Ala | Phe | Leu | Ile | Phe | Leu | |
| | | | | 245 | | | | | 250 | | | | | 255 | | |
| Leu | Leu | Gly | Ile | Cys | Trp | Cys | Gln | Cys | Cys | Pro | His | Thr | Cys | Cys | Cys | |
| | | | 260 | | | | | 265 | | | | | 270 | | | |
| Tyr | Val | Arg | Cys | Pro | Cys | Cys | Pro | Asp | Lys | Cys | Cys | Cys | Pro | Glu | Ala | |
| | | 275 | | | | | 280 | | | | | 285 | | | | |
| Leu | Tyr | Ala | Ala | Gly | Lys | Ala | Ala | Thr | Ser | Gly | Val | Pro | Ser | Ile | Tyr | |
| | 290 | | | | | 295 | | | | | 300 | | | | | |
| Ala | Pro | Ser | Thr | Tyr | Ala | His | Leu | Ser | Pro | Ala | Lys | Thr | Pro | Pro | Pro | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | |
| Pro | Ala | Met | Ile | Pro | Met | Gly | Pro | Ala | Tyr | Asn | Gly | Tyr | Pro | Gly | Gly | |
| | | | | 325 | | | | | 330 | | | | | 335 | | |
| Tyr | Pro | Gly | Asp | Val | Asp | Arg | Ser | Ser | Ser | Ala | Gly | Gly | Gln | Gly | Ser | |
| | | | 340 | | | | | 345 | | | | | 350 | | | |
| Tyr | Val | Pro | Leu | Leu | Arg | Asp | Thr | Asp | Ser | Ser | Val | Ala | Ser | Glu | Val | |
| | | 355 | | | | | 360 | | | | | 365 | | | | |
| Arg | Ser | Gly | Tyr | Arg | Ile | Gln | Ala | Ser | Gln | Gln | Asp | Ser | Ser | Met | Arg | |
| | 370 | | | | | 375 | | | | | 380 | | | | | |
| Val | Leu | Tyr | Tyr | Met | Glu | Lys | Glu | Leu | Ala | Asn | Phe | Asp | Pro | Ser | Arg | |
| 385 | | | | | 390 | | | | | 395 | | | | | 400 | |
| Pro | Gly | Pro | Pro | Ser | Gly | Arg | Val | Glu | Arg | Ala | Met | Ser | Glu | Val | Thr | |
| | | | | 405 | | | | | 410 | | | | | 415 | | |
| Ser | Leu | His | Glu | Asp | Asp | Trp | Arg | Ser | Arg | Pro | Ser | Arg | Gly | Pro | Ala | |
| | | | 420 | | | | | 425 | | | | | 430 | | | |
| Leu | Thr | Pro | Ile | Arg | Asp | Glu | Glu | Trp | Gly | Gly | His | Ser | Pro | Arg | Ser | |
| | | 435 | | | | | 440 | | | | | 445 | | | | |

Glu Glu Ala Tyr Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp
595 600 605
Ser Gln Ala Ser Arg Glu Arg Arg Leu Lys Lys Asn Leu Ala Leu Ser
610 615 620
Arg Glu Ser Leu Val Val
625 630

<210> 6
<211> 1954
<212> DNA
<213> Homo sapiens

<220>
<221> allele
<222> 595
<223> 9-3-324 : polymorphic base C or T

<220>
<221> allele
<222> 987
<223> 9-7-325 : polymorphic base A or G

<220>
<221> allele
<222> 1158
<223> 9-9-246 : polymorphic base G or C

<220>
<221> allele
<222> 1454
<223> LSRX9f13-BM : polymorphic base deletion of AGG

<220>
<221> allele
<222> 1875
<223> LSRX9f14-BM : polymorphic base T or G

<400> 6
tggagtgtgg ctcggaggac cgcggcggtt caagcacctt tctcccccat atctgaaagc 60
atgccctttg tccacgtcgt ttacgtcat taaaacttcc aga atg caa cag gac 115
Met Gln Gln Asp
1
gga ctt gga gta ggg aca agg aac gga agt ggg aag ggg agg agc gtg 163
Gly Leu Gly Val Gly Thr Arg Asn Gly Ser Gly Lys Gly Arg Ser Val
5 10 15 20
cac ccc tcc tgg cct tgg tgc gcg ccg cgc ccc cta agg tac ttt gga 211
His Pro Ser Trp Pro Trp Cys Ala Pro Arg Pro Leu Arg Tyr Phe Gly
25 30 35
agg gac gcg cgg gcc aga cgc gcc cag acg gcc gcg atg gcg ctg ttg 259
Arg Asp Ala Arg Ala Arg Arg Ala Gln Thr Ala Ala Met Ala Leu Leu
40 45 50
gcc ggc ggg ctc tcc aga ggg ctg ggc tcc cac ccg gcc gcc gca ggc 307
Ala Gly Gly Leu Ser Arg Gly Leu Gly Ser His Pro Ala Ala Ala Gly
55 60 65
cgg gac gcg gtc gtc ttc gtg tgg ctt ctg ctt agc acc tgg tgc aca 355
Arg Asp Ala Val Val Phe Val Trp Leu Leu Leu Ser Thr Trp Cys Thr
70 75 80
gct cct gcc agg gcc atc cag gtg acc gtg tcc aac ccc tac cac gtg 403
Ala Pro Ala Arg Ala Ile Gln Val Thr Val Ser Asn Pro Tyr His Val

| | | | | |
|---|-----|-----|-----|------|
| 85 | 90 | 95 | 100 | |
| gtg atc ctc ttc cag cct gtg acc ctg ccc tgt acc tac cag atg acc | | | | 451 |
| Val Ile Leu Phe Gln Pro Val Thr Leu Pro Cys Thr Tyr Gln Met Thr | | | | |
| | 105 | 110 | 115 | |
| tcg acc ccc acg caa ccc atc gtc atc tgg aag tac aag tct ttc tgc | | | | 499 |
| Ser Thr Pro Thr Gln Pro Ile Val Ile Trp Lys Tyr Lys Ser Phe Cys | | | | |
| | 120 | 125 | 130 | |
| cgg gac cgc atc gcc gat gcc ttc tcc ccg gcc agc gtc gac aac cag | | | | 547 |
| Arg Asp Arg Ile Ala Asp Ala Phe Ser Pro Ala Ser Val Asp Asn Gln | | | | |
| | 135 | 140 | 145 | |
| ctc aat gcc cag ctg gca gcc ggg aac cca ggc tac aac ccc tac gty | | | | 595 |
| Leu Asn Ala Gln Leu Ala Ala Gly Asn Pro Gly Tyr Asn Pro Tyr Val | | | | |
| | 150 | 155 | 160 | |
| gag tgc cag gac agc gtg cgc acc gtc agg gtc gtg gcc acc aag cag | | | | 643 |
| Glu Cys Gln Asp Ser Val Arg Thr Val Arg Val Val Ala Thr Lys Gln | | | | |
| | 165 | 170 | 175 | 180 |
| ggc aac gct gtg acc ctg gga gat tac tac cag ggc cgg agg att acc | | | | 691 |
| Gly Asn Ala Val Thr Leu Gly Asp Tyr Tyr Gln Gly Arg Arg Ile Thr | | | | |
| | 185 | 190 | 195 | |
| atc acc gga aat gct gac ctg acc ttt gac cag acg gcg tgg ggg gac | | | | 739 |
| Ile Thr Gly Asn Ala Asp Leu Thr Phe Asp Gln Thr Ala Trp Gly Asp | | | | |
| | 200 | 205 | 210 | |
| agt ggt gtg tat tac tgc tcc gtg gtc tca gcc cag gac ctc cag ggg | | | | 787 |
| Ser Gly Val Tyr Tyr Cys Ser Val Val Ser Ala Gln Asp Leu Gln Gly | | | | |
| | 215 | 220 | 225 | |
| aac aat gag gcc tac gca gag ctc atc gtc ctt gtg tat gcc gcc ggc | | | | 835 |
| Asn Asn Glu Ala Tyr Ala Glu Leu Ile Val Leu Val Tyr Ala Ala Gly | | | | |
| | 230 | 235 | 240 | |
| aaa gca gcc acc tca ggt gtt ccc agc att tat gcc ccc agc acc tat | | | | 883 |
| Lys Ala Ala Thr Ser Gly Val Pro Ser Ile Tyr Ala Pro Ser Thr Tyr | | | | |
| | 245 | 250 | 255 | 260 |
| gcc cac ctg tct ccc gcc aag acc cca ccc cca cca gct atg att ccc | | | | 931 |
| Ala His Leu Ser Pro Ala Lys Thr Pro Pro Pro Ala Met Ile Pro | | | | |
| | 265 | 270 | 275 | |
| atg ggc cct gcc tac aac ggg tac cct gga gga tac cct gga gac gtt | | | | 979 |
| Met Gly Pro Ala Tyr Asn Gly Tyr Pro Gly Gly Tyr Pro Gly Asp Val | | | | |
| | 280 | 285 | 290 | |
| gac agg art agc tca gct ggt ggc caa ggc tcc tat gta ccc ctg ctt | | | | 1027 |
| Asp Arg Xaa Ser Ser Ala Gly Gly Gln Gly Ser Tyr Val Pro Leu Leu | | | | |
| | 295 | 300 | 305 | |
| cgg gac acg gac agc agt gtg gcc tct gaa gtc cgc agt ggc tac agg | | | | 1075 |
| Arg Asp Thr Asp Ser Ser Val Ala Ser Glu Val Arg Ser Gly Tyr Arg | | | | |
| | 310 | 315 | 320 | |
| att cag gcc agc cag cag gac gac tcc atg cgg gtc ctg tac tac atg | | | | 1123 |
| Ile Gln Ala Ser Gln Gln Asp Asp Ser Met Arg Val Leu Tyr Tyr Met | | | | |
| | 325 | 330 | 335 | 340 |
| gag aag gag ctg gcc aac ttc gac cct tct cga cst ggc ccc ccc agt | | | | 1171 |
| Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser Arg Xaa Gly Pro Pro Ser | | | | |
| | 345 | 350 | 355 | |
| ggc cgt gtg gag cgg gcc atg agt gaa gtc acc tcc ctc cac gag gac | | | | 1219 |
| Gly Arg Val Glu Arg Ala Met Ser Glu Val Thr Ser Leu His Glu Asp | | | | |
| | 360 | 365 | 370 | |
| gac tgg cga tct cgg cct tcc cgg ggc cct gcc ctc acc ccg atc cgg | | | | 1267 |
| Asp Trp Arg Ser Arg Pro Ser Arg Gly Pro Ala Leu Thr Pro Ile Arg | | | | |
| | 375 | 380 | 385 | |
| gat gag gag tgg ggt ggc cac tcc ccc cgg agt ccc agg gga tgg gac | | | | 1315 |
| Asp Glu Glu Trp Gly Gly His Ser Pro Arg Ser Pro Arg Gly Trp Asp | | | | |
| | 390 | 395 | 400 | |
| cag gag ccc gcc agg gag cag gca ggc ggg ggc tgg cgg gcc agg cgg | | | | 1363 |

| | | |
|---|---|---------|
| Gln Glu Pro Ala Arg | Glu Gln Ala Gly Gly Gly Trp Arg Ala Arg Arg | |
| 405 | 410 | 415 420 |
| ccc cgg gcc cgc tcc gtg gac gcc ctg gac gac ctc acc ccg ccg agc | | 1411 |
| Pro Arg Ala Arg Ser Val Asp Ala Leu Asp Asp Leu Thr Pro Pro Ser | | |
| | 425 430 435 | |
| acc gcc gag tca ggg agc agg tct ccc acg agt aat ggt ggg aga agc | | 1459 |
| Thr Ala Glu Ser Gly Ser Arg Ser Pro Thr Ser Asn Gly Gly Arg Ser | | |
| | 440 445 450 | |
| cgg gcc tac atg ccc ccg cgg agc cgc agc cgg gac gac ctc tat gac | | 1507 |
| Arg Ala Tyr Met Pro Pro Arg Ser Arg Ser Arg Asp Asp Leu Tyr Asp | | |
| | 455 460 465 | |
| caa gac gac tgc agg gac ttc cca cgc tcc cgg gac ccc cac tac gac | | 1555 |
| Gln Asp Asp Ser Arg Asp Phe Pro Arg Ser Arg Asp Pro His Tyr Asp | | |
| | 470 475 480 | |
| gac ttc agg tct cgg gag cgc cct cct gcc gac ccc agg tcc cac cac | | 1603 |
| Asp Phe Arg Ser Arg Glu Arg Pro Pro Ala Asp Pro Arg Ser His His | | |
| | 485 490 495 500 | |
| cac cgt acc cgg gac cct cgg gac aac ggc tcc agg tcc ggg gac ctc | | 1651 |
| His Arg Thr Arg Asp Pro Arg Asp Asn Gly Ser Arg Ser Gly Asp Leu | | |
| | 505 510 515 | |
| ccc tat gat ggg cgg cta ctg gag gag gct gtg agg aag aag ggg tcg | | 1699 |
| Pro Tyr Asp Gly Arg Leu Leu Glu Glu Ala Val Arg Lys Lys Gly Ser | | |
| | 520 525 530 | |
| gag gag agg agg aga ccc cac aag gag gag gag gaa gag gcc tac tac | | 1747 |
| Glu Glu Arg Arg Arg Pro His Lys Glu Glu Glu Glu Glu Ala Tyr Tyr | | |
| | 535 540 545 | |
| ccg ccc gcg ccg ccc ccg tac tcg gag acc gac tcg cag gcg tcc cga | | 1795 |
| Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp Ser Gln Ala Ser Arg | | |
| | 550 555 560 | |
| gag cgc agg ctc aag aag aac ttg gcc ctg agt cgg gaa agt tta gtc | | 1843 |
| Glu Arg Arg Leu Lys Lys Asn Leu Ala Leu Ser Arg Glu Ser Leu Val | | |
| | 565 570 575 580 | |
| gtc tga tctgacgttt tctacgtagc ttttgkatTT ttttttttaa tttgaaggaa | | 1899 |
| Val * | | |
| cactgatgaa gccctgccat acccctcccg agtctaataa aacgtataat cacia | | 1954 |

<210> 7

<211> 581

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> 295

<223> 9-7-325 : polymorphic amino acid Ser or Asn

<220>

<221> VARIANT

<222> 352

<223> 9-9-246 : polymorphic amino acid Pro or Arg

<220>

<221> VARIANT

<222> 451

<223> LSRX9f13-BM : polymorphic amino acid deletion of Arg

<400> 7

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gln | Gln | Asp | Gly | Leu | Gly | Val | Gly | Thr | Arg | Asn | Gly | Ser | Gly | Lys |
| 1 | | | | | 5 | | | 10 | | | | | | 15 | |

| | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | | | 485 | | | | | | 490 | | | | | | 495 |
| Arg | Ser | His | His | His | Arg | Thr | Arg | Asp | Pro | Arg | Asp | Asn | Gly | Ser | Arg | |
| | | | 500 | | | | | 505 | | | | | 510 | | | |
| Ser | Gly | Asp | Leu | Pro | Tyr | Asp | Gly | Arg | Leu | Leu | Glu | Glu | Ala | Val | Arg | |
| | | 515 | | | | | 520 | | | | | 525 | | | | |
| Lys | Lys | Gly | Ser | Glu | Glu | Arg | Arg | Arg | Pro | His | Lys | Glu | Glu | Glu | Glu | |
| | 530 | | | | | 535 | | | | | 540 | | | | | |
| Glu | Ala | Tyr | Tyr | Pro | Pro | Ala | Pro | Pro | Pro | Tyr | Ser | Glu | Thr | Asp | Ser | |
| 545 | | | | | 550 | | | | | 555 | | | | | 560 | |
| Gln | Ala | Ser | Arg | Glu | Arg | Arg | Leu | Lys | Lys | Asn | Leu | Ala | Leu | Ser | Arg | |
| | | | | 565 | | | | | 570 | | | | | 575 | | |
| Glu | Ser | Leu | Val | Val | | | | | | | | | | | | |
| | | | 580 | | | | | | | | | | | | | |

<210> 8

<211> 2097

<212> DNA

<213> Rattus norvegicus

<400> 8

| | | | | | | | |
|---|------------|------------|------------|-------------------------|-------------|-----|-----|
| accgctcacc | aggtcagttg | tccccggaaa | gccgaaggca | tgagcttcgc | ccaagttctt | | 60 |
| tttatgggtt | agaactcctc | cagagcgggg | gaaaaaggac | ttggaatagg | ggcggggacgg | | 120 |
| agcacgcacc | cttctccgcc | ttggttctcg | ccgcgcccc | tactctcggg | atacttggga | | 180 |
| ggggacgcgc | gggcaccgtc | gctgctagac | ggccgcg | atg gcg ccg gcg gcc ggc | | | 235 |
| | | | Met | Ala | Pro | Ala | Gly |
| | | | 1 | | | 5 | |
| gcg tgt gct ggg gcg cct gac tcc cac cca gct acc gtg gtc ttc gtg | | | | | | | 283 |
| Ala Cys Ala Gly Ala Pro Asp Ser His Pro Ala Thr Val Val Phe Val | | | | | | | |
| | 10 | | 15 | | 20 | | |
| tgt ctc ttt ctc atc att ttc tgc cca gac cct gcc agt gcc atc cag | | | | | | | 331 |
| Cys Leu Phe Leu Ile Ile Phe Cys Pro Asp Pro Ala Ser Ala Ile Gln | | | | | | | |
| | 25 | | 30 | | 35 | | |
| gtg act gtg tct gac ccc tac cac gta gtg atc ctg ttc cag cca gtg | | | | | | | 379 |
| Val Thr Val Ser Asp Pro Tyr His Val Val Ile Leu Phe Gln Pro Val | | | | | | | |
| | 40 | | 45 | | 50 | | |
| acc ctg ccc tgc acc tat cag atg agc aac act ctc aca gtc ccc atc | | | | | | | 427 |
| Thr Leu Pro Cys Thr Tyr Gln Met Ser Asn Thr Leu Thr Val Pro Ile | | | | | | | |
| | 55 | | 60 | | 65 | | 70 |
| gtg atc tgg aag tac aag tca ttc tgc cgg gac cgt att gcc gat gcc | | | | | | | 475 |
| Val Ile Trp Lys Tyr Lys Ser Phe Cys Arg Asp Arg Ile Ala Asp Ala | | | | | | | |
| | 75 | | 80 | | 85 | | |
| ttc tct cct gcc agt gtg gac aac cag cta aat gcc cag ttg gca gct | | | | | | | 523 |
| Phe Ser Pro Ala Ser Val Asp Asn Gln Leu Asn Ala Gln Leu Ala Ala | | | | | | | |
| | 90 | | 95 | | 100 | | |
| ggc aac ccc ggc tac aac ccc tat gtg gag tgc cag gac agt gta cgc | | | | | | | 571 |
| Gly Asn Pro Gly Tyr Asn Pro Tyr Val Glu Cys Gln Asp Ser Val Arg | | | | | | | |
| | 105 | | 110 | | 115 | | |
| act gtc agg gtg gtg gcc acc aaa cag ggc aat gcg gtg acc ctg gga | | | | | | | 619 |
| Thr Val Arg Val Val Ala Thr Lys Gln Gly Asn Ala Val Thr Leu Gly | | | | | | | |
| | 120 | | 125 | | 130 | | |
| gac tac tac caa ggc agg agg atc acc ata aca gga aat gct gac ctg | | | | | | | 667 |
| Asp Tyr Tyr Gln Gly Arg Arg Ile Thr Ile Thr Gly Asn Ala Asp Leu | | | | | | | |
| | 135 | | 140 | | 145 | | 150 |
| acc ttc gag cag aca gcc tgg gga gac agt gga gtg tat tac tgc tct | | | | | | | 715 |
| Thr Phe Glu Gln Thr Ala Trp Gly Asp Ser Gly Val Tyr Tyr Cys Ser | | | | | | | |
| | 155 | | 160 | | 165 | | |
| gtg gtc tcg gcc caa gat ctg gat gga aac aac gag gcg tac gca gag | | | | | | | 763 |
| Val Val Ser Ala Gln Asp Leu Asp Gly Asn Asn Glu Ala Tyr Ala Glu | | | | | | | |
| | 170 | | 175 | | 180 | | |

| | |
|---|------|
| ctc atc gtc ctt ggc agg acc tca gag gcc cct gag ctc cta cct ggt | 811 |
| Leu Ile Val Leu Gly Arg Thr Ser Glu Ala Pro Glu Leu Leu Pro Gly | |
| 185 190 195 | |
| ttt cgg gcg ggg ccc ttg gaa gat tgg ctc ttt gtg gtc gtg gtc tgc | 859 |
| Phe Arg Ala Gly Pro Leu Glu Asp Trp Leu Phe Val Val Val Cys | |
| 200 205 210 | |
| ctg gcg agc ctc ctc ctc ttc ctc ctc ctg ggc atc tgc tgg tgc cag | 907 |
| Leu Ala Ser Leu Leu Leu Phe Leu Leu Leu Gly Ile Cys Trp Cys Gln | |
| 215 220 225 230 | |
| tgc tgt cct cac acc tgc tgc tgc tat gtc cga tgt ccc tgc tgc cca | 955 |
| Cys Cys Pro His Thr Cys Cys Cys Tyr Val Arg Cys Pro Cys Cys Pro | |
| 235 240 245 | |
| gac aag tgc tgt tgc cct gag gct ctt tat gct gct ggc aaa gca gcc | 1003 |
| Asp Lys Cys Cys Cys Pro Glu Ala Leu Tyr Ala Ala Gly Lys Ala Ala | |
| 250 255 260 | |
| acc tca ggt gtc ccg agc atc tat gcc ccc agc atc tat acc cac ctc | 1051 |
| Thr Ser Gly Val Pro Ser Ile Tyr Ala Pro Ser Ile Tyr Thr His Leu | |
| 265 270 275 | |
| tca cct gcc aag acc cca cca cct ccg cct gcc atg att ccc atg ggc | 1099 |
| Ser Pro Ala Lys Thr Pro Pro Pro Pro Pro Ala Met Ile Pro Met Gly | |
| 280 285 290 | |
| cct ccc tat ggg tac cct gga gac ttt gac aga cat agc tca gtt ggt | 1147 |
| Pro Pro Tyr Gly Tyr Pro Gly Asp Phe Asp Arg His Ser Ser Val Gly | |
| 295 300 305 310 | |
| ggc cac agc tcc caa gta ccc ctg ctg cgt gac gtg gat ggc agt gta | 1195 |
| Gly His Ser Ser Gln Val Pro Leu Leu Arg Asp Val Asp Gly Ser Val | |
| 315 320 325 | |
| tct tca gaa gta cga agt ggc tac agg atc cag gct aac cag caa gat | 1243 |
| Ser Ser Glu Val Arg Ser Gly Tyr Arg Ile Gln Ala Asn Gln Gln Asp | |
| 330 335 340 | |
| gac tcc atg agg gtc cta tac tat atg gag aaa gag cta gcc aac ttt | 1291 |
| Asp Ser Met Arg Val Leu Tyr Tyr Met Glu Lys Glu Leu Ala Asn Phe | |
| 345 350 355 | |
| gac cct tcc cga cct ggc cct ccc aat ggc aga gtg gaa cgg gcc atg | 1339 |
| Asp Pro Ser Arg Pro Gly Pro Pro Asn Gly Arg Val Glu Arg Ala Met | |
| 360 365 370 | |
| agt gaa gta acc tcc ctc cat gaa gat gac tgg cga tcg agg cct tcc | 1387 |
| Ser Glu Val Thr Ser Leu His Glu Asp Asp Trp Arg Ser Arg Pro Ser | |
| 375 380 385 390 | |
| agg gct cct gcc ctc acc ccc atc agg gat gag gag tgg aat cgc cac | 1435 |
| Arg Ala Pro Ala Leu Thr Pro Ile Arg Asp Glu Glu Trp Asn Arg His | |
| 395 400 405 | |
| tcc cca cag agt ccc aga aca tgg gag cag gaa ccc ctt caa gaa caa | 1483 |
| Ser Pro Gln Ser Pro Arg Thr Trp Glu Gln Glu Pro Leu Gln Glu Gln | |
| 410 415 420 | |
| cca agg ggt ggt tgg ggg tct gga cgc cct cgg gcc cgc tct gtg gat | 1531 |
| Pro Arg Gly Gly Trp Gly Ser Gly Arg Pro Arg Ala Arg Ser Val Asp | |
| 425 430 435 | |
| gct cta gat gat atc aac cgg cct ggc tcc act gaa tca gga cgg tct | 1579 |
| Ala Leu Asp Asp Ile Asn Arg Pro Gly Ser Thr Glu Ser Gly Arg Ser | |
| 440 445 450 | |
| tct ccc cca agt agt gga cgg aga gga cgg gcc tat gca cct cca aga | 1627 |
| Ser Pro Pro Ser Ser Gly Arg Arg Gly Arg Ala Tyr Ala Pro Pro Arg | |
| 455 460 465 470 | |
| agt cgc agc cgg gat gac ctc tat gac ccg gac gat cct agg gac ttg | 1675 |
| Ser Arg Ser Arg Asp Asp Leu Tyr Asp Pro Asp Asp Pro Arg Asp Leu | |
| 475 480 485 | |
| cca cat tcc cga gat ccc cac tat tat gac gac atc agg tct aga gat | 1723 |
| Pro His Ser Arg Asp Pro His Tyr Tyr Asp Asp Ile Arg Ser Arg Asp | |

| | | | | | | |
|---|-----|--|-----|--|-----|------|
| | 490 | | 495 | | 500 | |
| cca cgt gct gac ccc aga tcc cgt cag cga tcc cga gat cct cgg gat | | | | | | 1771 |
| Pro Arg Ala Asp Pro Arg Ser Arg Gln Arg Ser Arg Asp Pro Arg Asp | | | | | | |
| | 505 | | 510 | | 515 | |
| gct ggc ttc agg tca agg gac cct cag tat gat ggg cga cta tta gaa | | | | | | 1819 |
| Ala Gly Phe Arg Ser Arg Asp Pro Gln Tyr Asp Gly Arg Leu Leu Glu | | | | | | |
| | 520 | | 525 | | 530 | |
| gag gct tta aag aaa aag ggg tcg ggc gag aga agg agg gtt tac agg | | | | | | 1867 |
| Glu Ala Leu Lys Lys Lys Gly Ser Gly Glu Arg Arg Arg Val Tyr Arg | | | | | | |
| | 535 | | 540 | | 545 | 550 |
| gag gaa gaa gag gaa gag gag ggc caa tac ccc cca gca cct cca cct | | | | | | 1915 |
| Glu Glu Glu Glu Glu Glu Glu Gly Gln Tyr Pro Pro Ala Pro Pro Pro | | | | | | |
| | 555 | | 560 | | 565 | |
| tac tca gag act gac tcg cag gcc tca cgg gag agg agg ctg aaa aag | | | | | | 1963 |
| Tyr Ser Glu Thr Asp Ser Gln Ala Ser Arg Glu Arg Arg Leu Lys Lys | | | | | | |
| | 570 | | 575 | | 580 | |
| aat ttg gcc ctg agt cgg gaa agt tta gtc gtc tga tccacgtttt | | | | | | 2009 |
| Asn Leu Ala Leu Ser Arg Glu Ser Leu Val Val * | | | | | | |
| | 585 | | 590 | | | |
| gtatgtagct tttgtacttt ttttttaatt ggaatcaata ttgatgaaac ttcaagccta | | | | | | 2069 |
| ataaaatgtc taatcacaaa aaaaaaaa | | | | | | 2097 |

<210> 9
 <211> 593
 <212> PRT
 <213> Rattus norvegicus

<400> 9
 Met Ala Pro Ala Ala Gly Ala Cys Ala Gly Ala Pro Asp Ser His Pro
 1 5 10 15
 Ala Thr Val Val Phe Val Cys Leu Phe Leu Ile Ile Phe Cys Pro Asp
 20 25 30
 Pro Ala Ser Ala Ile Gln Val Thr Val Ser Asp Pro Tyr His Val Val
 35 40 45
 Ile Leu Phe Gln Pro Val Thr Leu Pro Cys Thr Tyr Gln Met Ser Asn
 50 55 60
 Thr Leu Thr Val Pro Ile Val Ile Trp Lys Tyr Lys Ser Phe Cys Arg
 65 70 75 80
 Asp Arg Ile Ala Asp Ala Phe Ser Pro Ala Ser Val Asp Asn Gln Leu
 85 90 95
 Asn Ala Gln Leu Ala Ala Gly Asn Pro Gly Tyr Asn Pro Tyr Val Glu
 100 105 110
 Cys Gln Asp Ser Val Arg Thr Val Arg Val Val Ala Thr Lys Gln Gly
 115 120 125
 Asn Ala Val Thr Leu Gly Asp Tyr Tyr Gln Gly Arg Arg Ile Thr Ile
 130 135 140
 Thr Gly Asn Ala Asp Leu Thr Phe Glu Gln Thr Ala Trp Gly Asp Ser
 145 150 155 160
 Gly Val Tyr Tyr Cys Ser Val Val Ser Ala Gln Asp Leu Asp Gly Asn
 165 170 175
 Asn Glu Ala Tyr Ala Glu Leu Ile Val Leu Gly Arg Thr Ser Glu Ala
 180 185 190
 Pro Glu Leu Leu Pro Gly Phe Arg Ala Gly Pro Leu Glu Asp Trp Leu
 195 200 205
 Phe Val Val Val Val Cys Leu Ala Ser Leu Leu Leu Phe Leu Leu Leu
 210 215 220
 Gly Ile Cys Trp Cys Gln Cys Cys Pro His Thr Cys Cys Cys Tyr Val
 225 230 235 240
 Arg Cys Pro Cys Cys Pro Asp Lys Cys Cys Cys Pro Glu Ala Leu Tyr

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|------|
| | | | 10 | | | | 15 | | | | 20 | | | | | | |
| tgt | ctc | ttt | ctc | atc | att | ttc | tgc | cca | gac | cct | gcc | agt | gcc | atc | cag | | 331 |
| Cys | Leu | Phe | Leu | Ile | Ile | Phe | Cys | Pro | Asp | Pro | Ala | Ser | Ala | Ile | Gln | | |
| | | 25 | | | | | 30 | | | | 35 | | | | | | |
| gtg | act | gtg | tct | gac | ccc | tac | cac | gta | gtg | atc | ctg | ttc | cag | cca | gtg | | 379 |
| Val | Thr | Val | Ser | Asp | Pro | Tyr | His | Val | Val | Ile | Leu | Phe | Gln | Pro | Val | | |
| | 40 | | | | | 45 | | | | | 50 | | | | | | |
| acc | ctg | ccc | tgc | acc | tat | cag | atg | agc | aac | act | ctc | aca | gtc | ccc | atc | | 427 |
| Thr | Leu | Pro | Cys | Thr | Tyr | Gln | Met | Ser | Asn | Thr | Leu | Thr | Val | Pro | Ile | | |
| | 55 | | | | 60 | | | | 65 | | | | 70 | | | | |
| gtg | atc | tgg | aag | tac | aag | tca | ttc | tgc | cgg | gac | cgt | att | gcc | gat | gcc | | 475 |
| Val | Ile | Trp | Lys | Tyr | Lys | Ser | Phe | Cys | Arg | Asp | Arg | Ile | Ala | Asp | Ala | | |
| | | | 75 | | | | | | 80 | | | | 85 | | | | |
| ttc | tct | cct | gcc | agt | gtg | gac | aac | cag | cta | aat | gcc | cag | ttg | gca | gct | | 523 |
| Phe | Ser | Pro | Ala | Ser | Val | Asp | Asn | Gln | Leu | Asn | Ala | Gln | Leu | Ala | Ala | | |
| | | 90 | | | | | 95 | | | | 100 | | | | | | |
| ggc | aac | ccc | ggc | tac | aac | ccc | tat | gtg | gag | tgc | cag | gac | agt | gta | cgc | | 571 |
| Gly | Asn | Pro | Gly | Tyr | Asn | Pro | Tyr | Val | Glu | Cys | Gln | Asp | Ser | Val | Arg | | |
| | 105 | | | | | 110 | | | | | 115 | | | | | | |
| act | gtc | agg | gtg | gtg | gcc | acc | aaa | cag | ggc | aat | gcg | gtg | acc | ctg | gga | | 619 |
| Thr | Val | Arg | Val | Val | Ala | Thr | Lys | Gln | Gly | Asn | Ala | Val | Thr | Leu | Gly | | |
| | 120 | | | | | 125 | | | | | 130 | | | | | | |
| gac | tac | tac | caa | ggc | agg | agg | atc | acc | ata | aca | gga | aat | gct | gac | ctg | | 667 |
| Asp | Tyr | Tyr | Gln | Gly | Arg | Arg | Ile | Thr | Ile | Thr | Gly | Asn | Ala | Asp | Leu | | |
| | 135 | | | | 140 | | | | | 145 | | | | 150 | | | |
| acc | ttc | gag | cag | aca | gcc | tgg | gga | gac | agt | gga | gtg | tat | tac | tgc | tct | | 715 |
| Thr | Phe | Glu | Gln | Thr | Ala | Trp | Gly | Asp | Ser | Gly | Val | Tyr | Tyr | Cys | Ser | | |
| | | | 155 | | | | | 160 | | | | | 165 | | | | |
| gtg | gtc | tcg | gcc | caa | gat | ctg | gat | gga | aac | aac | gag | gcg | tac | gca | gag | | 763 |
| Val | Val | Ser | Ala | Gln | Asp | Leu | Asp | Gly | Asn | Asn | Glu | Ala | Tyr | Ala | Glu | | |
| | | 170 | | | | | 175 | | | | | 180 | | | | | |
| ctc | atc | gtc | ctt | gat | tgg | ctc | ttt | gtg | gtc | gtg | gtc | tgc | ctg | gcg | agc | | 811 |
| Leu | Ile | Val | Leu | Asp | Trp | Leu | Phe | Val | Val | Val | Val | Cys | Leu | Ala | Ser | | |
| | | 185 | | | | 190 | | | | | | 195 | | | | | |
| ctc | ctc | ctc | ttc | ctc | ctc | ctg | ggc | atc | tgc | tgg | tgc | cag | tgc | tgt | cct | | 859 |
| Leu | Leu | Leu | Phe | Leu | Leu | Leu | Gly | Ile | Cys | Trp | Cys | Gln | Cys | Cys | Pro | | |
| | 200 | | | | | 205 | | | | | 210 | | | | | | |
| cac | acc | tgc | tgc | tgc | tat | gtc | cga | tgt | ccc | tgc | tgc | cca | gac | aag | tgc | | 907 |
| His | Thr | Cys | Cys | Cys | Tyr | Val | Arg | Cys | Pro | Cys | Cys | Pro | Asp | Lys | Cys | | |
| | 215 | | | | 220 | | | | 225 | | | | | 230 | | | |
| tgt | tgc | cct | gag | gct | ctt | tat | gct | gct | ggc | aaa | gca | gcc | acc | tca | ggt | | 955 |
| Cys | Cys | Pro | Glu | Ala | Leu | Tyr | Ala | Ala | Gly | Lys | Ala | Ala | Thr | Ser | Gly | | |
| | | | 235 | | | | | 240 | | | | | 245 | | | | |
| gtc | ccg | agc | atc | tat | gcc | ccc | agc | atc | tat | acc | cac | ctc | tca | cct | gcc | | 1003 |
| Val | Pro | Ser | Ile | Tyr | Ala | Pro | Ser | Ile | Tyr | Thr | His | Leu | Ser | Pro | Ala | | |
| | | 250 | | | | | | 255 | | | | 260 | | | | | |
| aag | acc | cca | cca | cct | ccg | cct | gcc | atg | att | ccc | atg | ggc | cct | ccc | tat | | 1051 |
| Lys | Thr | Pro | Pro | Pro | Pro | Pro | Ala | Met | Ile | Pro | Met | Gly | Pro | Pro | Tyr | | |
| | 265 | | | | | 270 | | | | 275 | | | | | | | |
| ggg | tac | cct | gga | gac | ttt | gac | aga | cat | agc | tca | gtt | ggt | ggc | cac | agc | | 1099 |
| Gly | Tyr | Pro | Gly | Asp | Phe | Asp | Arg | His | Ser | Ser | Val | Gly | Gly | His | Ser | | |
| | 280 | | | | 285 | | | | | 290 | | | | | | | |
| tcc | caa | gta | ccc | ctg | ctg | cgt | gac | gtg | gat | ggc | agt | gta | tct | tca | gaa | | 1147 |
| Ser | Gln | Val | Pro | Leu | Leu | Arg | Asp | Val | Asp | Gly | Ser | Val | Ser | Ser | Glu | | |
| | 295 | | | | 300 | | | | 305 | | | | | 310 | | | |
| gta | cga | agt | ggc | tac | agg | atc | cag | gct | aac | cag | caa | gat | gac | tcc | atg | | 1195 |
| Val | Arg | Ser | Gly | Tyr | Arg | Ile | Gln | Ala | Asn | Gln | Gln | Asp | Asp | Ser | Met | | |
| | | | 315 | | | | | 320 | | | | 325 | | | | | |
| agg | gtc | cta | tac | tat | atg | gag | aaa | gag | cta | gcc | aac | ttt | gac | cct | tcc | | 1243 |

| | |
|---|------|
| Arg Val Leu Tyr Tyr Met Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser | |
| 330 335 340 | |
| cga cct ggc cct ccc aat ggc aga gtg gaa cgg gcc atg agt gaa gta | 1291 |
| Arg Pro Gly Pro Pro Asn Gly Arg Val Glu Arg Ala Met Ser Glu Val | |
| 345 350 355 | |
| acc tcc ctc cat gaa gat gac tgg cga tcg agg cct tcc agg gct cct | 1339 |
| Thr Ser Leu His Glu Asp Asp Trp Arg Ser Arg Pro Ser Arg Ala Pro | |
| 360 365 370 | |
| gcc ctc acc ccc atc agg gat gag gag tgg aat cgc cac tcc cca cag | 1387 |
| Ala Leu Thr Pro Ile Arg Asp Glu Glu Trp Asn Arg His Ser Pro Gln | |
| 375 380 385 390 | |
| agt ccc aga aca tgg gag cag gaa ccc ctt caa gaa caa cca agg ggt | 1435 |
| Ser Pro Arg Thr Trp Glu Gln Glu Pro Leu Gln Glu Gln Pro Arg Gly | |
| 395 400 405 | |
| ggt tgg ggg tct gga cgc cct cgg gcc cgc tct gtg gat gct cta gat | 1483 |
| Gly Trp Gly Ser Gly Arg Pro Arg Ala Arg Ser Val Asp Ala Leu Asp | |
| 410 415 420 | |
| gat atc aac cgg cct ggc tcc act gaa tca gga cgg tct tct ccc cca | 1531 |
| Asp Ile Asn Arg Pro Gly Ser Thr Glu Ser Gly Arg Ser Ser Pro Pro | |
| 425 430 435 | |
| agt agt gga cgg aga gga cgg gcc tat gca cct cca aga agt cgc agc | 1579 |
| Ser Ser Gly Arg Arg Gly Arg Ala Tyr Ala Pro Pro Arg Ser Arg Ser | |
| 440 445 450 | |
| cgg gat gac ctc tat gac ccg gac gat cct agg gac ttg cca cat tcc | 1627 |
| Arg Asp Asp Leu Tyr Asp Pro Asp Asp Pro Arg Asp Leu Pro His Ser | |
| 455 460 465 470 | |
| cga gat ccc cac tat tat gac gac atc agg tct aga gat cca cgt gct | 1675 |
| Arg Asp Pro His Tyr Tyr Asp Asp Ile Arg Ser Arg Asp Pro Arg Ala | |
| 475 480 485 | |
| gac ccc aga tcc cgt cag cga tcc cga gat cct cgg gat gct ggc ttc | 1723 |
| Asp Pro Arg Ser Arg Gln Arg Ser Arg Asp Pro Arg Asp Ala Gly Phe | |
| 490 495 500 | |
| agg tca agg gac cct cag tat gat ggg cga cta tta gaa gag gct tta | 1771 |
| Arg Ser Arg Asp Pro Gln Tyr Asp Gly Arg Leu Leu Glu Glu Ala Leu | |
| 505 510 515 | |
| aag aaa aag ggg tcg ggc gag aga agg agg gtt tac agg gag gaa gaa | 1819 |
| Lys Lys Lys Gly Ser Gly Glu Arg Arg Arg Val Tyr Arg Glu Glu Glu | |
| 520 525 530 | |
| gag gaa gag gag ggc caa tac ccc cca gca cct cca cct tac tca gag | 1867 |
| Glu Glu Glu Glu Gly Gln Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu | |
| 535 540 545 550 | |
| act gac tcg cag gcc tca cgg gag agg agg ctg aaa aag aat ttg gcc | 1915 |
| Thr Asp Ser Gln Ala Ser Arg Glu Arg Arg Leu Lys Lys Asn Leu Ala | |
| 555 560 565 | |
| ctg agt cgg gaa agt tta gtc gtc tga tccacgtttt gtatgtagct | 1962 |
| Leu Ser Arg Glu Ser Leu Val Val * | |
| 570 575 | |
| tttgtacttt ttttttaatt ggaatcaata ttgatgaaac ttcaagccta ataaaatgtc | 2022 |
| taatcacaaa aaaaaaaaa | 2040 |

<210> 11

<211> 574

<212> PRT

<213> Rattus norvegicus

<400> 11

| | |
|---|--|
| Met Ala Pro Ala Ala Gly Ala Cys Ala Gly Ala Pro Asp Ser His Pro | |
| 1 5 10 15 | |
| Ala Thr Val Val Phe Val Cys Leu Phe Leu Ile Ile Phe Cys Pro Asp | |

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|--|
| | | | 20 | | | | | 25 | | | | | | | 30 | | |
| Pro | Ala | Ser | Ala. | Ile | Gln | Val | Thr | Val | Ser | Asp | Pro | Tyr | His | Val | Val | | |
| | | 35 | | | | | 40 | | | | | 45 | | | | | |
| Ile | Leu | Phe | Gln | Pro | Val | Thr | Leu | Pro | Cys | Thr | Tyr | Gln | Met | Ser | Asn | | |
| | 50 | | | | | 55 | | | | | 60 | | | | | | |
| Thr | Leu | Thr | Val | Pro | Ile | Val | Ile | Trp | Lys | Tyr | Lys | Ser | Phe | Cys | Arg | | |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 | | |
| Asp | Arg | Ile | Ala | Asp | Ala | Phe | Ser | Pro | Ala | Ser | Val | Asp | Asn | Gln | Leu | | |
| | | | | 85 | | | | | 90 | | | | | 95 | | | |
| Asn | Ala | Gln | Leu | Ala | Ala | Gly | Asn | Pro | Gly | Tyr | Asn | Pro | Tyr | Val | Glu | | |
| | | | 100 | | | | | 105 | | | | | 110 | | | | |
| Cys | Gln | Asp | Ser | Val | Arg | Thr | Val | Arg | Val | Val | Ala | Thr | Lys | Gln | Gly | | |
| | 115 | | | | | | 120 | | | | | 125 | | | | | |
| Asn | Ala | Val | Thr | Leu | Gly | Asp | Tyr | Tyr | Gln | Gly | Arg | Arg | Ile | Thr | Ile | | |
| | 130 | | | | | 135 | | | | | 140 | | | | | | |
| Thr | Gly | Asn | Ala | Asp | Leu | Thr | Phe | Glu | Gln | Thr | Ala | Trp | Gly | Asp | Ser | | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | | |
| Gly | Val | Tyr | Tyr | Cys | Ser | Val | Val | Ser | Ala | Gln | Asp | Leu | Asp | Gly | Asn | | |
| | | | | 165 | | | | | 170 | | | | | 175 | | | |
| Asn | Glu | Ala | Tyr | Ala | Glu | Leu | Ile | Val | Leu | Asp | Trp | Leu | Phe | Val | Val | | |
| | | | 180 | | | | | 185 | | | | | 190 | | | | |
| Val | Val | Cys | Leu | Ala | Ser | Leu | Leu | Leu | Phe | Leu | Leu | Leu | Gly | Ile | Cys | | |
| | | 195 | | | | | 200 | | | | | 205 | | | | | |
| Trp | Cys | Gln | Cys | Cys | Pro | His | Thr | Cys | Cys | Cys | Tyr | Val | Arg | Cys | Pro | | |
| | 210 | | | | | 215 | | | | | 220 | | | | | | |
| Cys | Cys | Pro | Asp | Lys | Cys | Cys | Cys | Pro | Glu | Ala | Leu | Tyr | Ala | Ala | Gly | | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | | |
| Lys | Ala | Ala | Thr | Ser | Gly | Val | Pro | Ser | Ile | Tyr | Ala | Pro | Ser | Ile | Tyr | | |
| | | | | 245 | | | | | 250 | | | | | 255 | | | |
| Thr | His | Leu | Ser | Pro | Ala | Lys | Thr | Pro | Pro | Pro | Pro | Pro | Ala | Met | Ile | | |
| | | | 260 | | | | | 265 | | | | | 270 | | | | |
| Pro | Met | Gly | Pro | Pro | Tyr | Gly | Tyr | Pro | Gly | Asp | Phe | Asp | Arg | His | Ser | | |
| | | 275 | | | | | 280 | | | | | 285 | | | | | |
| Ser | Val | Gly | Gly | His | Ser | Ser | Gln | Val | Pro | Leu | Leu | Arg | Asp | Val | Asp | | |
| | 290 | | | | | 295 | | | | | 300 | | | | | | |
| Gly | Ser | Val | Ser | Ser | Glu | Val | Arg | Ser | Gly | Tyr | Arg | Ile | Gln | Ala | Asn | | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | | |
| Gln | Gln | Asp | Asp | Ser | Met | Arg | Val | Leu | Tyr | Tyr | Met | Glu | Lys | Glu | Leu | | |
| | | | | 325 | | | | | 330 | | | | | 335 | | | |
| Ala | Asn | Phe | Asp | Pro | Ser | Arg | Pro | Gly | Pro | Pro | Asn | Gly | Arg | Val | Glu | | |
| | | | 340 | | | | | 345 | | | | | | | | | |

Pro Arg Asp Ala Gly Phe Arg Ser Arg Asp Pro Gln Tyr Asp Gly Arg
500 505 510
Leu Leu Glu Glu Ala Leu Lys Lys Lys Gly Ser Gly Glu Arg Arg Arg
515 520 525
Val Tyr Arg Glu Glu Glu Glu Glu Glu Gly Gln Tyr Pro Pro Ala
530 535 540
Pro Pro Pro Tyr Ser Glu Thr Asp Ser Gln Ala Ser Arg Glu Arg Arg
545 550 555 560
Leu Lys Lys Asn Leu Ala Leu Ser Arg Glu Ser Leu Val Val
565 570

<210> 12
<211> 1893
<212> DNA
<213> Rattus norvegicus

<400> 12
accgctcacc aggtcagttg tccccggaaa gccgaaggca tgagcttcgc ccaagttctt 60
tttatgggtt agaactcctc cagagcgggg gaaaaaggac ttggaatagg ggcgggacgg 120
agcacgcacc cttctcgcgc ttggttctcg ccgcgcccc tactctcggg atacttgggga 180
ggggacgcgc gggcaccgtc gctgctagac ggccgcg atg gcg ccg gcg gcc ggc 235
Met Ala Pro Ala Ala Gly
1 5
gcg tgt gct ggg gcg cct gac tcc cac cca gct acc gtg gtc ttc gtg 283
Ala Cys Ala Gly Ala Pro Asp Ser His Pro Ala Thr Val Val Phe Val
10 15 20
tgt ctc ttt ctc atc att ttc tgc cca gac cct gcc agt gcc atc cag 331
Cys Leu Phe Leu Ile Ile Phe Cys Pro Asp Pro Ala Ser Ala Ile Gln
25 30 35
gtg act gtg tct gac ccc tac cac gta gtg atc ctg ttc cag cca gtg 379
Val Thr Val Ser Asp Pro Tyr His Val Val Ile Leu Phe Gln Pro Val
40 45 50
acc ctg ccc tgc acc tat cag atg agc aac act ctc aca gtc ccc atc 427
Thr Leu Pro Cys Thr Tyr Gln Met Ser Asn Thr Leu Thr Val Pro Ile
55 60 65 70
gtg atc tgg aag tac aag tca ttc tgc cgg gac cgt att gcc gat gcc 475
Val Ile Trp Lys Tyr Lys Ser Phe Cys Arg Asp Arg Ile Ala Asp Ala
75 80 85
ttc tct cct gcc agt gtg gac aac cag cta aat gcc cag ttg gca gct 523
Phe Ser Pro Ala Ser Val Asp Asn Gln Leu Asn Ala Gln Leu Ala Ala
90 95 100
ggc aac ccc ggc tac aac ccc tat gtg gag tgc cag gac agt gta cgc 571
Gly Asn Pro Gly Tyr Asn Pro Tyr Val Glu Cys Gln Asp Ser Val Arg
105 110 115
act gtc agg gtg gtg gcc acc aaa cag ggc aat gcg gtg acc ctg gga 619
Thr Val Arg Val Val Ala Thr Lys Gln Gly Asn Ala Val Thr Leu Gly
120 125 130
gac tac tac caa ggc agg agg atc acc ata aca gga aat gct gac ctg 667
Asp Tyr Tyr Gln Gly Arg Arg Ile Thr Ile Thr Gly Asn Ala Asp Leu
135 140 145 150
acc ttc gag cag aca gcc tgg gga gac agt gga gtg tat tac tgc tct 715
Thr Phe Glu Gln Thr Ala Trp Gly Asp Ser Gly Val Tyr Tyr Cys Ser
155 160 165
gtg gtc tcg gcc caa gat ctg gat gga aac aac gag gcg tac gca gag 763
Val Val Ser Ala Gln Asp Leu Asp Gly Asn Asn Glu Ala Tyr Ala Glu
170 175 180
ctc atc gtc ctt gtt tat gct gct ggc aaa gca gcc acc tca ggt gtc 811
Leu Ile Val Leu Val Tyr Ala Ala Gly Lys Ala Ala Thr Ser Gly Val
185 190 195

| | |
|---|------|
| ccg agc atc tat gcc ccc agc atc tat acc cac ctc tca cct gcc aag | 859 |
| Pro Ser Ile Tyr Ala Pro Ser Ile Tyr Thr His Leu Ser Pro Ala Lys | |
| 200 205 210 | |
| acc cca cca cct ccg cct gcc atg att ccc atg ggc cct ccc tat ggg | 907 |
| Thr Pro Pro Pro Pro Ala Met Ile Pro Met Gly Pro Pro Tyr Gly | |
| 215 220 225 230 | |
| tac cct gga gac ttt gac aga cat agc tca gtt ggt ggc cac agc tcc | 955 |
| Tyr Pro Gly Asp Phe Asp Arg His Ser Ser Val Gly Gly His Ser Ser | |
| 235 240 245 | |
| caa gta ccc ctg ctg cgt gac gtg gat ggc agt gta tct tca gaa gta | 1003 |
| Gln Val Pro Leu Leu Arg Asp Val Asp Gly Ser Val Ser Ser Glu Val | |
| 250 255 260 | |
| cga agt ggc tac agg atc cag gct aac cag caa gat gac tcc atg agg | 1051 |
| Arg Ser Gly Tyr Arg Ile Gln Ala Asn Gln Gln Asp Asp Ser Met Arg | |
| 265 270 275 | |
| gtc cta tac tat atg gag aaa gag cta gcc aac ttt gac cct tcc cga | 1099 |
| Val Leu Tyr Tyr Met Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser Arg | |
| 280 285 290 | |
| cct ggc cct ccc aat ggc aga gtg gaa cgg gcc atg agt gaa gta acc | 1147 |
| Pro Gly Pro Pro Asn Gly Arg Val Glu Arg Ala Met Ser Glu Val Thr | |
| 295 300 305 310 | |
| tcc ctc cat gaa gat gac tgg cga tcg agg cct tcc agg gct cct gcc | 1195 |
| Ser Leu His Glu Asp Trp Arg Ser Arg Pro Ser Arg Ala Pro Ala | |
| 315 320 325 | |
| ctc acc ccc atc agg gat gag gag tgg aat cgc cac tcc cca cag agt | 1243 |
| Leu Thr Pro Ile Arg Asp Glu Glu Trp Asn Arg His Ser Pro Gln Ser | |
| 330 335 340 | |
| ccc aga aca tgg gag cag gaa ccc ctt caa gaa caa cca agg ggt ggt | 1291 |
| Pro Arg Thr Trp Glu Gln Glu Pro Leu Gln Glu Gln Pro Arg Gly Gly | |
| 345 350 355 | |
| tgg ggg tct gga cgc cct cgg gcc cgc tct gtg gat gct cta gat gat | 1339 |
| Trp Gly Ser Gly Arg Pro Arg Ala Arg Ser Val Asp Ala Leu Asp Asp | |
| 360 365 370 | |
| atc aac cgg cct ggc tcc act gaa tca gga cgg tct tct ccc cca agt | 1387 |
| Ile Asn Arg Pro Gly Ser Thr Glu Ser Gly Arg Ser Ser Pro Pro Ser | |
| 375 380 385 390 | |
| agt gga cgg aga gga cgg gcc tat gca cct cca aga agt cgc agc cgg | 1435 |
| Ser Gly Arg Arg Gly Arg Ala Tyr Ala Pro Pro Arg Ser Arg Ser Arg | |
| 395 400 405 | |
| gat gac ctc tat gac ccg gac gat cct agg gac ttg cca cat tcc cga | 1483 |
| Asp Asp Leu Tyr Asp Pro Asp Asp Pro Arg Asp Leu Pro His Ser Arg | |
| 410 415 420 | |
| gat ccc cac tat tat gac gac atc agg tct aga gat cca cgt gct gac | 1531 |
| Asp Pro His Tyr Tyr Asp Asp Ile Arg Ser Arg Asp Pro Arg Ala Asp | |
| 425 430 435 | |
| ccc aga tcc cgt cag cga tcc cga gat cct cgg gat gct ggc ttc agg | 1579 |
| Pro Arg Ser Arg Gln Arg Ser Arg Asp Pro Arg Asp Ala Gly Phe Arg | |
| 440 445 450 | |
| tca agg gac cct cag tat gat ggg cga cta tta gaa gag gct tta aag | 1627 |
| Ser Arg Asp Pro Gln Tyr Asp Gly Arg Leu Leu Glu Glu Ala Leu Lys | |
| 455 460 465 470 | |
| aaa aag ggg tcg ggc gag aga agg agg gtt tac agg gag gaa gaa gag | 1675 |
| Lys Lys Gly Ser Gly Glu Arg Arg Arg Val Tyr Arg Glu Glu Glu Glu | |
| 475 480 485 | |
| gaa gag gag ggc caa tac ccc cca gca cct cca cct tac tca gag act | 1723 |
| Glu Glu Glu Gly Gln Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr | |
| 490 495 500 | |
| gac tcg cag gcc tca cgg gag agg agg ctg aaa aag aat ttg gcc ctg | 1771 |
| Asp Ser Gln Ala Ser Arg Glu Arg Arg Leu Lys Lys Asn Leu Ala Leu | |

Val Asp Ala Leu Asp Asp Ile Asn Arg Pro Gly Ser Thr Glu Ser Gly
 370 375 380
 Arg Ser Ser Pro Pro Ser Ser Gly Arg Arg Gly Arg Ala Tyr Ala Pro
 385 390 395 400
 Pro Arg Ser Arg Ser Arg Asp Asp Leu Tyr Asp Pro Asp Asp Pro Arg
 405 410 415
 Asp Leu Pro His Ser Arg Asp Pro His Tyr Tyr Asp Asp Ile Arg Ser
 420 425 430
 Arg Asp Pro Arg Ala Asp Pro Arg Ser Arg Gln Arg Ser Arg Asp Pro
 435 440 445
 Arg Asp Ala Gly Phe Arg Ser Arg Asp Pro Gln Tyr Asp Gly Arg Leu
 450 455 460
 Leu Glu Glu Ala Leu Lys Lys Lys Gly Ser Gly Glu Arg Arg Arg Val
 465 470 475 480
 Tyr Arg Glu Glu Glu Glu Glu Glu Gly Gln Tyr Pro Pro Ala Pro
 485 490 495
 Pro Pro Tyr Ser Glu Thr Asp Ser Gln Ala Ser Arg Glu Arg Arg Leu
 500 505 510
 Lys Lys Asn Leu Ala Leu Ser Arg Glu Ser Leu Val Val
 515 520 525

<210> 14

<211> 1886

<212> DNA

<213> Mus musculus

<400> 14

gcaccgtcgc tgctagacgg ccgcg atg gcg ccg gcg gcc agc gcg tgt gct 52
 Met Ala Pro Ala Ala Ser Ala Cys Ala
 1 5
 ggg gcg cct ggc tcc cac ccg gcc acc acg atc ttc gtg tgt ctt ttt 100
 Gly Ala Pro Gly Ser His Pro Ala Thr Thr Ile Phe Val Cys Leu Phe
 10 15 20 25
 ctc atc att tac tgc cca gac cgt gcc agt gcc atc cag gtg acc gtg 148
 Leu Ile Ile Tyr Cys Pro Asp Arg Ala Ser Ala Ile Gln Val Thr Val
 30 35 40
 cct gac ccc tac cac gta gtg atc ctg ttc cag cca gtg aca cta cac 196
 Pro Asp Pro Tyr His Val Val Ile Leu Phe Gln Pro Val Thr Leu His
 45 50 55
 tgc acc tac cag atg agc aat acc ctc aca gcc cct atc gtg atc tgg 244
 Cys Thr Tyr Gln Met Ser Asn Thr Leu Thr Ala Pro Ile Val Ile Trp
 60 65 70
 aag tat aag tcg ttc tgt cgg gac cgt gtt gcc gac gcc ttc tcc cct 292
 Lys Tyr Lys Ser Phe Cys Arg Asp Arg Val Ala Asp Ala Phe Ser Pro
 75 80 85
 gcc agc gtg gac aac cag ctc aac gcc cag ctg gcg gct ggc aac ccc 340
 Ala Ser Val Asp Asn Gln Leu Asn Ala Gln Leu Ala Ala Gly Asn Pro
 90 95 100 105
 ggc tac aac ccc tat gtg gag tgc cag gac agc gta cgc act gtc agg 388
 Gly Tyr Asn Pro Tyr Val Glu Cys Gln Asp Ser Val Arg Thr Val Arg
 110 115 120
 gtg gtg gcc acc aaa cag ggc aat gct gtg acc ctg gga gac tac tac 436
 Val Val Ala Thr Lys Gln Gly Asn Ala Val Thr Leu Gly Asp Tyr Tyr
 125 130 135
 cag ggc agg aga atc acc atc aca gga aat gct ggc ctg acc ttc gag 484
 Gln Gly Arg Arg Ile Thr Ile Thr Gly Asn Ala Gly Leu Thr Phe Glu
 140 145 150
 cag acg gcc tgg gga gac agt gga gtg tat tac tgc tcc gtg gtc tca 532
 Gln Thr Ala Trp Gly Asp Ser Gly Val Tyr Tyr Cys Ser Val Val Ser

| | | | |
|---|-----|-----|------|
| 155 | 160 | 165 | |
| gcc caa gat ctg gat ggg aac aac gag gcg tac gca gag ctc att gtc | | | 580 |
| Ala Gln Asp Leu Asp Gly Asn Asn Glu Ala Tyr Ala Glu Leu Ile Val | | | |
| 170 | 175 | 180 | 185 |
| ctt ggc agg acc tca gaa gcc cct gag ctc cta cct ggt ttt cgg gcg | | | 628 |
| Leu Gly Arg Thr Ser Glu Ala Pro Glu Leu Leu Pro Gly Phe Arg Ala | | | |
| | 190 | 195 | 200 |
| ggg ccc ttg gaa gat tgg ctc ttt gtg gtc gtg gtc tgc ctg gca agc | | | 676 |
| Gly Pro Leu Glu Asp Trp Leu Phe Val Val Val Val Cys Leu Ala Ser | | | |
| | 205 | 210 | 215 |
| ctc ctc ttc ttc ctc ctc ctg ggc atc tgc tgg tgc cag tgc tgt ccc | | | 724 |
| Leu Leu Phe Phe Leu Leu Leu Gly Ile Cys Trp Cys Gln Cys Cys Pro | | | |
| | 220 | 225 | 230 |
| cac acc tgc tgc tgc tat gtc aga tgt ccc tgc tgc cca gac aag tgc | | | 772 |
| His Thr Cys Cys Cys Tyr Val Arg Cys Pro Cys Cys Pro Asp Lys Cys | | | |
| | 235 | 240 | 245 |
| tgt tgc cct gag gcc ctt tat gct gct ggc aaa gca gcc acc tca ggt | | | 820 |
| Cys Cys Pro Glu Ala Leu Tyr Ala Ala Gly Lys Ala Ala Thr Ser Gly | | | |
| | 250 | 255 | 260 |
| gtg cca agc atc tat gcc ccc agc atc tat acc cac ctc tct cct gcc | | | 868 |
| Val Pro Ser Ile Tyr Ala Pro Ser Ile Tyr Thr His Leu Ser Pro Ala | | | |
| | 270 | 275 | 280 |
| aag act ccg cca cct ccg cct gcc atg att ccc atg cgt cct ccc tat | | | 916 |
| Lys Thr Pro Pro Pro Pro Ala Met Ile Pro Met Arg Pro Pro Tyr | | | |
| | 285 | 290 | 295 |
| ggg tac cct gga gac ttt gac agg acc agc tca gtt ggt ggc cac agc | | | 964 |
| Gly Tyr Pro Gly Asp Phe Asp Arg Thr Ser Ser Val Gly Gly His Ser | | | |
| | 300 | 305 | 310 |
| tcc cag gtg ccc ctg ctg cgt gaa gtg gat ggg agc gta tct tca gaa | | | 1012 |
| Ser Gln Val Pro Leu Leu Arg Glu Val Asp Gly Ser Val Ser Ser Glu | | | |
| | 315 | 320 | 325 |
| gta cga agt ggc tac agg atc cag gct aac cag caa gat gac tcc atg | | | 1060 |
| Val Arg Ser Gly Tyr Arg Ile Gln Ala Asn Gln Asp Asp Ser Met | | | |
| | 330 | 335 | 340 |
| agg gtc cta tac tat atg gag aag gag cta gcc aac ttc gat cct tcc | | | 1108 |
| Arg Val Leu Tyr Tyr Met Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser | | | |
| | 350 | 355 | 360 |
| cgg cct ggc cct ccc aat ggc cga gtg gaa cgg gcc atg agt gaa gta | | | 1156 |
| Arg Pro Gly Pro Pro Asn Gly Arg Val Glu Arg Ala Met Ser Glu Val | | | |
| | 365 | 370 | 375 |
| acc tcc ctc cat gaa gat gac tgg cga tct cgg cct tcc agg gct cct | | | 1204 |
| Thr Ser Leu His Glu Asp Asp Trp Arg Ser Arg Pro Ser Arg Ala Pro | | | |
| | 380 | 385 | 390 |
| gcc ctc aca ccc atc agg gat gag gag tgg aat cgc cac tcc cct cgg | | | 1252 |
| Ala Leu Thr Pro Ile Arg Asp Glu Glu Trp Asn Arg His Ser Pro Arg | | | |
| | 395 | 400 | 405 |
| agt ccc aga aca tgg gag cag gaa ccc ctt caa gaa cag cca agg ggt | | | 1300 |
| Ser Pro Arg Thr Trp Glu Gln Glu Pro Leu Gln Glu Gln Pro Arg Gly | | | |
| | 410 | 415 | 420 |
| ggg tgg ggg tct ggg cgg cct cgg gcc cgc tct gtg gat gct cta gat | | | 1348 |
| Gly Trp Gly Ser Gly Arg Pro Arg Ala Arg Ser Val Asp Ala Leu Asp | | | |
| | 430 | 435 | 440 |
| gac atc aac cgg cct ggc tcc act gaa tca gga agg tct tct ccc cca | | | 1396 |
| Asp Ile Asn Arg Pro Gly Ser Thr Glu Ser Gly Arg Ser Ser Pro Pro | | | |
| | 445 | 450 | 455 |
| agt agt gga cgg aga ggg cgg gcc tat gca cct ccg aga agt cgc agc | | | 1444 |
| Ser Ser Gly Arg Arg Gly Arg Ala Tyr Ala Pro Pro Arg Ser Arg Ser | | | |
| | 460 | 465 | 470 |
| cgg gat gac ctc tat gac ccc gac gat cct aga gac ttg cca cat tcc | | | 1492 |

| | |
|---|------|
| Arg Asp Asp Leu Tyr Asp Pro Asp Asp Pro Arg Asp Leu Pro His Ser | |
| 475 480 485 | |
| cga gat ccc cac tat tat gat gat ttg agg tct agg gat cca cgt gct | 1540 |
| Arg Asp Pro His Tyr Asp Asp Leu Arg Ser Arg Asp Pro Arg Ala | |
| 490 495 500 505 | |
| gac ccc aga tcc cgt cag cga tcc cac gat cct cgg gat gct ggc ttc | 1588 |
| Asp Pro Arg Ser Arg Gln Arg Ser His Asp Pro Arg Asp Ala Gly Phe | |
| 510 515 520 | |
| agg tca cgg gac cct cag tat gat ggg cga ctc tta gaa gag gct tta | 1636 |
| Arg Ser Arg Asp Pro Gln Tyr Asp Gly Arg Leu Leu Glu Glu Ala Leu | |
| 525 530 535 | |
| aag aaa aaa ggg gct ggg gag aga aga cgc gtt tac agg gag gaa gaa | 1684 |
| Lys Lys Lys Gly Ala Gly Glu Arg Arg Val Tyr Arg Glu Glu Glu | |
| 540 545 550 | |
| gaa gaa gaa gag gag ggc cac tat ccc cca gca cct ccg cct tac tct | 1732 |
| Glu Glu Glu Glu Glu Gly His Tyr Pro Pro Ala Pro Pro Pro Tyr Ser | |
| 555 560 565 | |
| gag act gac tcg cag gcc tcg agg gag cgg agg atg aaa aag aat ttg | 1780 |
| Glu Thr Asp Ser Gln Ala Ser Arg Glu Arg Arg Met Lys Lys Asn Leu | |
| 570 575 580 585 | |
| gcc ctg agt cgg gaa agt tta gtc gtc tga tccccacgttt tgttatgtag | 1830 |
| Ala Leu Ser Arg Glu Ser Leu Val Val * | |
| 590 595 | |
| cttttatact tttttaattg gaattatgat gaaactcttc accaagccta ataaaa | 1886 |
| <210> 15 | |
| <211> 1829 | |
| <212> DNA | |
| <213> Mus musculus | |
| <400> 15 | |
| gcaccgtcgc tgctagacgg ccgcg atg gcg ccg gcg gcc agc gcg tgt gct | 52 |
| Met Ala Pro Ala Ala Ser Ala Cys Ala | |
| 1 5 | |
| ggg gcg cct ggc tcc cac ccg gcc acc acg atc ttc gtg tgt ctt ttt | 100 |
| Gly Ala Pro Gly Ser His Pro Ala Thr Thr Ile Phe Val Cys Leu Phe | |
| 10 15 20 25 | |
| ctc atc att tac tgc cca gac cgt gcc agt gcc atc cag gtg acc gtg | 148 |
| Leu Ile Ile Tyr Cys Pro Asp Arg Ala Ser Ala Ile Gln Val Thr Val | |
| 30 35 40 | |
| cct gac ccc tac cac gta gtg atc ctg ttc cag cca gtg aca cta cac | 196 |
| Pro Asp Pro Tyr His Val Val Ile Leu Phe Gln Pro Val Thr Leu His | |
| 45 50 55 | |
| tgc acc tac cag atg agc aat acc ctc aca gcc cct atc gtg atc tgg | 244 |
| Cys Thr Tyr Gln Met Ser Asn Thr Leu Thr Ala Pro Ile Val Ile Trp | |
| 60 65 70 | |
| aag tat aag tcg ttc tgt cgg gac cgt gtt gcc gac gcc ttc tcc cct | 292 |
| Lys Tyr Lys Ser Phe Cys Arg Asp Arg Val Ala Asp Ala Phe Ser Pro | |
| 75 80 85 | |
| gcc agc gtg gac aac cag ctc aac gcc cag ctg gcg gct ggc aac ccc | 340 |
| Ala Ser Val Asp Asn Gln Leu Asn Ala Gln Leu Ala Ala Gly Asn Pro | |
| 90 95 100 105 | |
| ggc tac aac ccc tat gtg gag tgc cag gac agc gta cgc act gtc agg | 388 |
| Gly Tyr Asn Pro Tyr Val Glu Cys Gln Asp Ser Val Arg Thr Val Arg | |
| 110 115 120 | |
| gtg gtg gcc acc aaa cag ggc aat gct gtg acc ctg gga gac tac tac | 436 |
| Val Val Ala Thr Lys Gln Gly Asn Ala Val Thr Leu Gly Asp Tyr Tyr | |
| 125 130 135 | |
| cag ggc agg aga atc acc atc aca gga aat gct ggc ctg acc ttc gag | 484 |

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|--|
| Gln | Gly | Arg | Arg | Ile | Thr | Ile | Thr | Gly | Asn | Ala | Gly | Leu | Thr | Phe | Glu | | |
| | | 140 | | | | | 145 | | | | 150 | | | | | | |
| cag | acg | gcc | tgg | gga | gac | agt | gga | gtg | tat | tac | tgc | tcc | gtg | gtc | tca | 532 | |
| Gln | Thr | Ala | Trp | Gly | Asp | Ser | Gly | Val | Tyr | Tyr | Cys | Ser | Val | Val | Ser | | |
| | | 155 | | | | | 160 | | | | 165 | | | | | | |
| gcc | caa | gat | ctg | gat | ggg | aac | aac | gag | gcg | tac | gca | gag | ctc | att | gtc | 580 | |
| Ala | Gln | Asp | Leu | Asp | Gly | Asn | Asn | Glu | Ala | Tyr | Ala | Glu | Leu | Ile | Val | | |
| | | 170 | | | 175 | | | | | 180 | | | | | 185 | | |
| ctt | gat | tgg | ctc | ttt | gtg | gtc | gtg | gtc | tgc | ctg | gca | agc | ctc | ctc | ttc | 628 | |
| Leu | Asp | Trp | Leu | Phe | Val | Val | Val | Val | Cys | Leu | Ala | Ser | Leu | Leu | Phe | | |
| | | | | 190 | | | | | 195 | | | | | 200 | | | |
| ttc | ctc | ctc | ctg | ggc | atc | tgc | tgg | tgc | cag | tgc | tgt | ccc | cac | acc | tgc | 676 | |
| Phe | Leu | Leu | Leu | Gly | Ile | Cys | Trp | Cys | Gln | Cys | Cys | Pro | His | Thr | Cys | | |
| | | | 205 | | | | | 210 | | | | | 215 | | | | |
| tgc | tgc | tat | gtc | aga | tgt | ccc | tgc | tgc | cca | gac | aag | tgc | tgt | tgc | cct | 724 | |
| Cys | Cys | Tyr | Val | Arg | Cys | Pro | Cys | Cys | Pro | Asp | Lys | Cys | Cys | Cys | Pro | | |
| | | 220 | | | | | 225 | | | | 230 | | | | | | |
| gag | gcc | ctt | tat | gct | gct | ggc | aaa | gca | gcc | acc | tca | ggg | gtg | cca | agc | 772 | |
| Glu | Ala | Leu | Tyr | Ala | Ala | Gly | Lys | Ala | Ala | Thr | Ser | Gly | Val | Pro | Ser | | |
| | | 235 | | | | 240 | | | | 245 | | | | | | | |
| atc | tat | gcc | ccc | agc | atc | tat | acc | cac | ctc | tct | cct | gcc | aag | act | ccg | 820 | |
| Ile | Tyr | Ala | Pro | Ser | Ile | Tyr | Thr | His | Leu | Ser | Pro | Ala | Lys | Thr | Pro | | |
| | | 250 | | | 255 | | | | 260 | | | | | 265 | | | |
| cca | cct | ccg | cct | gcc | atg | att | ccc | atg | cgt | cct | ccc | tat | ggg | tac | cct | 868 | |
| Pro | Pro | Pro | Pro | Ala | Met | Ile | Pro | Met | Arg | Pro | Pro | Tyr | Gly | Tyr | Pro | | |
| | | | | 270 | | | | 275 | | | | | 280 | | | | |
| gga | gac | ttt | gac | agg | acc | agc | tca | gtt | ggg | ggc | cac | agc | tcc | cag | gtg | 916 | |
| Gly | Asp | Phe | Asp | Arg | Thr | Ser | Ser | Val | Gly | Gly | His | Ser | Ser | Gln | Val | | |
| | | 285 | | | | | 290 | | | | 295 | | | | | | |
| ccc | ctg | ctg | cgt | gaa | gtg | gat | ggg | agc | gta | tct | tca | gaa | gta | cga | agt | 964 | |
| Pro | Leu | Leu | Arg | Glu | Val | Asp | Gly | Ser | Val | Ser | Ser | Glu | Val | Arg | Ser | | |
| | | 300 | | | | | 305 | | | | 310 | | | | | | |
| ggc | tac | agg | atc | cag | gct | aac | cag | caa | gat | gac | tcc | atg | agg | gtc | cta | 1012 | |
| Gly | Tyr | Arg | Ile | Gln | Ala | Asn | Gln | Gln | Asp | Asp | Ser | Met | Arg | Val | Leu | | |
| | | 315 | | | | 320 | | | | 325 | | | | | | | |
| tac | tat | atg | gag | aag | gag | cta | gcc | aac | ttc | gat | cct | tcc | cgg | cct | ggc | 1060 | |
| Tyr | Tyr | Met | Glu | Lys | Glu | Leu | Ala | Asn | Phe | Asp | Pro | Ser | Arg | Pro | Gly | | |
| | | 330 | | | 335 | | | | 340 | | | | | 345 | | | |
| cct | ccc | aat | ggc | cga | gtg | gaa | cgg | gcc | atg | agt | gaa | gta | acc | tcc | ctc | 1108 | |
| Pro | Pro | Asn | Gly | Arg | Val | Glu | Arg | Ala | Met | Ser | Glu | Val | Thr | Ser | Leu | | |
| | | | 350 | | | | | 355 | | | | | 360 | | | | |
| cat | gaa | gat | gac | tgg | cga | tct | cgg | cct | tcc | agg | gct | cct | gcc | ctc | aca | 1156 | |
| His | Glu | Asp | Asp | Trp | Arg | Ser | Arg | Pro | Ser | Arg | Ala | Pro | Ala | Leu | Thr | | |
| | | 365 | | | | | 370 | | | | 375 | | | | | | |
| ccc | atc | agg | gat | gag | gag | tgg | aat | cgc | cac | tcc | cct | cgg | agt | ccc | aga | 1204 | |
| Pro | Ile | Arg | Asp | Glu | Glu | Trp | Asn | Arg | His | Ser | Pro | Arg | Ser | Pro | Arg | | |
| | | 380 | | | | | 385 | | | | 390 | | | | | | |
| aca | tgg | gag | cag | gaa | ccc | ctt | caa | gaa | cag | cca | agg | ggg | ggg | tgg | ggg | 1252 | |
| Thr | Trp | Glu | Gln | Glu | Pro | Leu | Gln | Glu | Gln | Pro | Arg | Gly | Gly | Trp | Gly | | |
| | | 395 | | | | 400 | | | | 405 | | | | | | | |
| tct | ggg | cgg | cct | cgg | gcc | cgc | tct | gtg | gat | gct | cta | gat | gac | atc | aac | 1300 | |
| Ser | Gly | Arg | Pro | Arg | Ala | Arg | Ser | Val | Asp | Ala | Leu | Asp | Asp | Ile | Asn | | |
| | | 410 | | | 415 | | | | 420 | | | | | 425 | | | |
| cgg | cct | ggc | tcc | act | gaa | tca | gga | agg | tct | tct | ccc | cca | agt | agt | gga | 1348 | |
| Arg | Pro | Gly | Ser | Thr | Glu | Ser | Gly | Arg | Ser | Ser | Pro | Pro | Ser | Ser | Gly | | |
| | | | 430 | | | | 435 | | | | | | 440 | | | | |
| cgg | aga | ggg | cgg | gcc | tat | gca | cct | ccg | aga | agt | cgc | agc | cgg | gat | gac | 1396 | |
| Arg | Arg | Gly | Arg | Ala | Tyr | Ala | Pro | Pro | Arg | Ser | Arg | Ser | Arg | Asp | Asp | | |
| | | 445 | | | | | 450 | | | | | | 455 | | | | |

| | |
|---|------|
| ctc tat gac ccc gac gat cct aga gac ttg cca cat tcc cga gat ccc | 1444 |
| Leu Tyr Asp Pro Asp Asp Pro Arg Asp Leu Pro His Ser Arg Asp Pro | |
| 460 465 470 | |
| cac tat tat gat gat ttg agg tct agg gat cca cgt gct gac ccc aga | 1492 |
| His Tyr Tyr Asp Asp Leu Arg Ser Arg Asp Pro Arg Ala Asp Pro Arg | |
| 475 480 485 | |
| tcc cgt cag cga tcc cac gat cct cgg gat gct ggc ttc agg tca cgg | 1540 |
| Ser Arg Gln Arg Ser His Asp Pro Arg Asp Ala Gly Phe Arg Ser Arg | |
| 490 495 500 505 | |
| gac cct cag tat gat ggg cga ctc tta gaa gag gct tta aag aaa aaa | 1588 |
| Asp Pro Gln Tyr Asp Gly Arg Leu Leu Glu Glu Ala Leu Lys Lys Lys | |
| 510 515 520 | |
| ggg gct ggg gag aga aga cgc gtt tac agg gag gaa gaa gaa gaa gaa | 1636 |
| Gly Ala Gly Glu Arg Arg Arg Val Tyr Arg Glu Glu Glu Glu Glu Glu | |
| 525 530 535 | |
| gag gag ggc cac tat ccc cca gca cct ccg cct tac tct gag act gac | 1684 |
| Glu Glu Gly His Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp | |
| 540 545 550 | |
| tcg cag gcc tcg agg gag cgg agg atg aaa aag aat ttg gcc ctg agt | 1732 |
| Ser Gln Ala Ser Arg Glu Arg Arg Met Lys Lys Asn Leu Ala Leu Ser | |
| 555 560 565 | |
| cgg gaa agt tta gtc gtc tga tccccagttt tgttatgtag cttttatact | 1783 |
| Arg Glu Ser Leu Val Val * | |
| 570 575 | |
| tttttaattg gaatattgat gaaactcttc accaagccta ataaaa | 1829 |
| <210> 16 | |
| <211> 1682 | |
| <212> DNA | |
| <213> Mus musculus | |
| <400> 16 | |
| gcaccgtcgc tgctagacgg ccgcg atg gcg ccg gcg gcc agc gcg tgt gct | 52 |
| Met Ala Pro Ala Ala Ser Ala Cys Ala | |
| 1 5 | |
| ggg gcg cct ggc tcc cac ccg gcc acc acg atc ttc gtg tgt ctt ttt | 100 |
| Gly Ala Pro Gly Ser His Pro Ala Thr Thr Ile Phe Val Cys Leu Phe | |
| 10 15 20 25 | |
| ctc atc att tac tgc cca gac cgt gcc agt gcc atc cag gtg acc gtg | 148 |
| Leu Ile Ile Tyr Cys Pro Asp Arg Ala Ser Ala Ile Gln Val Thr Val | |
| 30 35 40 | |
| cct gac ccc tac cac gta gtg atc ctg ttc cag cca gtg aca cta cac | 196 |
| Pro Asp Pro Tyr His Val Val Ile Leu Phe Gln Pro Val Thr Leu His | |
| 45 50 55 | |
| tgc acc tac cag atg agc aat acc ctc aca gcc cct atc gtg atc tgg | 244 |
| Cys Thr Tyr Gln Met Ser Asn Thr Leu Thr Ala Pro Ile Val Ile Trp | |
| 60 65 70 | |
| aag tat aag tcg ttc tgt cgg gac cgt gtt gcc gac gcc ttc tcc cct | 292 |
| Lys Tyr Lys Ser Phe Cys Arg Asp Arg Val Ala Asp Ala Phe Ser Pro | |
| 75 80 85 | |
| gcc agc gtg gac aac cag ctc aac gcc cag ctg gcg gct ggc aac ccc | 340 |
| Ala Ser Val Asp Asn Gln Leu Asn Ala Gln Leu Ala Ala Gly Asn Pro | |
| 90 95 100 105 | |
| ggc tac aac ccc tat gtg gag tgc cag gac agc gta cgc act gtc agg | 388 |
| Gly Tyr Asn Pro Tyr Val Glu Cys Gln Asp Ser Val Arg Thr Val Arg | |
| 110 115 120 | |
| gtg gtg gcc acc aaa cag ggc aat gct gtg acc ctg gga gac tac tac | 436 |
| Val Val Ala Thr Lys Gln Gly Asn Ala Val Thr Leu Gly Asp Tyr Tyr | |
| 125 130 135 | |

| | |
|---|------|
| cag ggc agg aga atc acc atc aca gga aat gct ggc ctg acc ttc gag | 484 |
| Gln Gly Arg Arg Ile Thr Ile Thr Gly Asn Ala Gly Leu Thr Phe Glu | |
| 140 145 150 | |
| cag acg gcc tgg gga gac agt gga gtg tat tac tgc tcc gtg gtc tca | 532 |
| Gln Thr Ala Trp Gly Asp Ser Gly Val Tyr Tyr Cys Ser Val Val Ser | |
| 155 160 165 | |
| gcc caa gat ctg gat ggg aac aac gag gcg tac gca gag ctc att gtc | 580 |
| Ala Gln Asp Leu Asp Gly Asn Asn Glu Ala Tyr Ala Glu Leu Ile Val | |
| 170 175 180 185 | |
| ctt gtt tat gct gct ggc aaa gca gcc acc tca ggt gtg cca agc atc | 628 |
| Leu Val Tyr Ala Ala Gly Lys Ala Ala Thr Ser Gly Val Pro Ser Ile | |
| 190 195 200 | |
| tat gcc ccc agc atc tat acc cac ctc tct cct gcc aag act ccg cca | 676 |
| Tyr Ala Pro Ser Ile Tyr Thr His Leu Ser Pro Ala Lys Thr Pro Pro | |
| 205 210 215 | |
| cct ccg cct gcc atg att ccc atg cgt cct ccc tat ggg tac cct gga | 724 |
| Pro Pro Pro Ala Met Ile Pro Met Arg Pro Pro Tyr Gly Tyr Pro Gly | |
| 220 225 230 | |
| gac ttt gac agg acc agc tca gtt ggt ggc cac agc tcc cag gtg ccc | 772 |
| Asp Phe Asp Arg Thr Ser Ser Val Gly Gly His Ser Ser Gln Val Pro | |
| 235 240 245 | |
| ctg ctg cgt gaa gtg gat ggg agc gta tct tca gaa gta cga agt ggc | 820 |
| Leu Leu Arg Glu Val Asp Gly Ser Val Ser Ser Glu Val Arg Ser Gly | |
| 250 255 260 265 | |
| tac agg atc cag gct aac cag caa gat gac tcc atg agg gtc cta tac | 868 |
| Tyr Arg Ile Gln Ala Asn Gln Gln Asp Asp Ser Met Arg Val Leu Tyr | |
| 270 275 280 | |
| tat atg gag aag gag cta gcc aac ttc gat cct tcc cgg cct ggc cct | 916 |
| Tyr Met Glu Lys Glu Leu Ala Asn Phe Asp Pro Ser Arg Pro Gly Pro | |
| 285 290 295 | |
| ccc aat ggc cga gtg gaa cgg gcc atg agt gaa gta acc tcc ctc cat | 964 |
| Pro Asn Gly Arg Val Glu Arg Ala Met Ser Glu Val Thr Ser Leu His | |
| 300 305 310 | |
| gaa gat gac tgg cga tct cgg cct tcc agg gct cct gcc ctc aca ccc | 1012 |
| Glu Asp Asp Trp Arg Ser Arg Pro Ser Arg Ala Pro Ala Leu Thr Pro | |
| 315 320 325 | |
| atc agg gat gag gag tgg aat cgc cac tcc cct cgg agt ccc aga aca | 1060 |
| Ile Arg Asp Glu Glu Trp Asn Arg His Ser Pro Arg Ser Pro Arg Thr | |
| 330 335 340 345 | |
| tgg gag cag gaa ccc ctt caa gaa cag cca agg ggt ggt tgg ggg tct | 1108 |
| Trp Glu Gln Glu Pro Leu Gln Glu Gln Pro Arg Gly Gly Trp Gly Ser | |
| 350 355 360 | |
| ggg cgg cct cgg gcc cgc tct gtg gat gct cta gat gac atc aac cgg | 1156 |
| Gly Arg Pro Arg Ala Arg Ser Val Asp Ala Leu Asp Asp Ile Asn Arg | |
| 365 370 375 | |
| cct ggc tcc act gaa tca gga agg tct tct ccc cca agt agt gga cgg | 1204 |
| Pro Gly Ser Thr Glu Ser Gly Arg Ser Ser Pro Pro Ser Ser Gly Arg | |
| 380 385 390 | |
| aga ggg cgg gcc tat gca cct ccg aga agt cgc agc cgg gat gac ctc | 1252 |
| Arg Gly Arg Ala Tyr Ala Pro Pro Arg Ser Arg Ser Arg Asp Asp Leu | |
| 395 400 405 | |
| tat gac ccc gac gat cct aga gac ttg cca cat tcc cga gat ccc cac | 1300 |
| Tyr Asp Pro Asp Asp Pro Arg Asp Leu Pro His Ser Arg Asp Pro His | |
| 410 415 420 425 | |
| tat tat gat gat ttg agg tct agg gat cca cgt gct gac ccc aga tcc | 1348 |
| Tyr Tyr Asp Asp Leu Arg Ser Arg Asp Pro Arg Ala Asp Pro Arg Ser | |
| 430 435 440 | |
| cgt cag cga tcc cac gat cct cgg gat gct ggc ttc agg tca cgg gac | 1396 |
| Arg Gln Arg Ser His Asp Pro Arg Asp Ala Gly Phe Arg Ser Arg Asp | |

| | | | | |
|---|-----|-----|-----|------|
| | 445 | 450 | 455 | |
| cct cag tat gat ggg cga ctc tta gaa gag gct tta aag aaa aaa ggg | | | | 1444 |
| Pro Gln Tyr Asp Gly Arg Leu Leu Glu Glu Ala Leu Lys Lys Lys Gly | | | | |
| | 460 | 465 | 470 | |
| gct ggg gag aga aga cgc gtt tac agg gag gaa gaa gaa gaa gag | | | | 1492 |
| Ala Gly Glu Arg Arg Arg Val Tyr Arg Glu Glu Glu Glu Glu Glu | | | | |
| | 475 | 480 | 485 | |
| gag ggc cac tat ccc cca gca cct ccg cct tac tct gag act gac tcg | | | | 1540 |
| Glu Gly His Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp Ser | | | | |
| | 490 | 495 | 500 | 505 |
| cag gcc tcg agg gag cgg agg atg aaa aag aat ttg gcc ctg agt cgg | | | | 1588 |
| Gln Ala Ser Arg Glu Arg Arg Met Lys Lys Asn Leu Ala Leu Ser Arg | | | | |
| | 510 | 515 | 520 | |
| gaa agt tta gtc gtc tga tcccacgttt tggtatgtag cttttatact | | | | 1636 |
| Glu Ser Leu Val Val * | | | | |
| | 525 | | | |
| tttttaattg gaatatgat gaaactcttc accaagccta ataaaa | | | | 1682 |

<210> 17
 <211> 594
 <212> PRT
 <213> Mus musculus

<400> 17

| | | | | |
|---|-----|-----|-----|--|
| Met Ala Pro Ala Ala Ser Ala Cys Ala Gly Ala Pro Gly Ser His Pro | | | | |
| 1 | 5 | 10 | 15 | |
| Ala Thr Thr Ile Phe Val Cys Leu Phe Leu Ile Ile Tyr Cys Pro Asp | | | | |
| | 20 | 25 | 30 | |
| Arg Ala Ser Ala Ile Gln Val Thr Val Pro Asp Pro Tyr His Val Val | | | | |
| | 35 | 40 | 45 | |
| Ile Leu Phe Gln Pro Val Thr Leu His Cys Thr Tyr Gln Met Ser Asn | | | | |
| | 50 | 55 | 60 | |
| Thr Leu Thr Ala Pro Ile Val Ile Trp Lys Tyr Lys Ser Phe Cys Arg | | | | |
| 65 | 70 | 75 | 80 | |
| Asp Arg Val Ala Asp Ala Phe Ser Pro Ala Ser Val Asp Asn Gln Leu | | | | |
| | 85 | 90 | 95 | |
| Asn Ala Gln Leu Ala Ala Gly Asn Pro Gly Tyr Asn Pro Tyr Val Glu | | | | |
| | 100 | 105 | 110 | |
| Cys Gln Asp Ser Val Arg Thr Val Arg Val Val Ala Thr Lys Gln Gly | | | | |
| | 115 | 120 | 125 | |
| Asn Ala Val Thr Leu Gly Asp Tyr Tyr Gln Gly Arg Arg Ile Thr Ile | | | | |
| | 130 | 135 | 140 | |
| Thr Gly Asn Ala Gly Leu Thr Phe Glu Gln Thr Ala Trp Gly Asp Ser | | | | |
| 145 | 150 | 155 | 160 | |
| Gly Val Tyr Tyr Cys Ser Val Val Ser Ala Gln Asp Leu Asp Gly Asn | | | | |
| | 165 | 170 | 175 | |
| Asn Glu Ala Tyr Ala Glu Leu Ile Val Leu Gly Arg Thr Ser Glu Ala | | | | |
| | 180 | 185 | 190 | |
| Pro Glu Leu Leu Pro Gly Phe Arg Ala Gly Pro Leu Glu Asp Trp Leu | | | | |
| | 195 | 200 | 205 | |
| Phe Val Val Val Val Cys Leu Ala Ser Leu Leu Phe Phe Leu Leu Leu | | | | |
| | 210 | 215 | 220 | |
| Gly Ile Cys Trp Cys Gln Cys Cys Pro His Thr Cys Cys Cys Tyr Val | | | | |
| 225 | 230 | 235 | 240 | |
| Arg Cys Pro Cys Cys Pro Asp Lys Cys Cys Cys Pro Glu Ala Leu Tyr | | | | |
| | 245 | 250 | 255 | |
| Ala Ala Gly Lys Ala Ala Thr Ser Gly Val Pro Ser Ile Tyr Ala Pro | | | | |
| | 260 | 265 | 270 | |
| Ser Ile Tyr Thr His Leu Ser Pro Ala Lys Thr Pro Pro Pro Pro Pro | | | | |

| | | |
|---|-----|-----|
| 275 | 280 | 285 |
| Ala Met Ile Pro Met Arg Pro Pro Tyr Gly Tyr Pro Gly Asp Phe Asp | | |
| 290 | 295 | 300 |
| Arg Thr Ser Ser Val Gly His Ser Ser Gln Val Pro Leu Leu Arg | | |
| 305 | 310 | 315 |
| Glu Val Asp Gly Ser Val Ser Ser Glu Val Arg Ser Gly Tyr Arg Ile | | |
| 325 | 330 | 335 |
| Gln Ala Asn Gln Gln Asp Asp Ser Met Arg Val Leu Tyr Tyr Met Glu | | |
| 340 | 345 | 350 |
| Lys Glu Leu Ala Asn Phe Asp Pro Ser Arg Pro Gly Pro Pro Asn Gly | | |
| 355 | 360 | 365 |
| Arg Val Glu Arg Ala Met Ser Glu Val Thr Ser Leu His Glu Asp Asp | | |
| 370 | 375 | 380 |
| Trp Arg Ser Arg Pro Ser Arg Ala Pro Ala Leu Thr Pro Ile Arg Asp | | |
| 385 | 390 | 395 |
| Glu Glu Trp Asn Arg His Ser Pro Arg Ser Pro Arg Thr Trp Glu Gln | | |
| 405 | 410 | 415 |
| Glu Pro Leu Gln Glu Gln Pro Arg Gly Gly Trp Gly Ser Gly Arg Pro | | |
| 420 | 425 | 430 |
| Arg Ala Arg Ser Val Asp Ala Leu Asp Asp Ile Asn Arg Pro Gly Ser | | |
| 435 | 440 | 445 |
| Thr Glu Ser Gly Arg Ser Ser Pro Pro Ser Ser Gly Arg Arg Gly Arg | | |
| 450 | 455 | 460 |
| Ala Tyr Ala Pro Pro Arg Ser Arg Ser Arg Asp Asp Leu Tyr Asp Pro | | |
| 465 | 470 | 475 |
| Asp Asp Pro Arg Asp Leu Pro His Ser Arg Asp Pro His Tyr Tyr Asp | | |
| 485 | 490 | 495 |
| Asp Leu Arg Ser Arg Asp Pro Arg Ala Asp Pro Arg Ser Arg Gln Arg | | |
| 500 | 505 | 510 |
| Ser His Asp Pro Arg Asp Ala Gly Phe Arg Ser Arg Asp Pro Gln Tyr | | |
| 515 | 520 | 525 |
| Asp Gly Arg Leu Leu Glu Glu Ala Leu Lys Lys Lys Gly Ala Gly Glu | | |
| 530 | 535 | 540 |
| Arg Arg Arg Val Tyr Arg Glu Glu Glu Glu Glu Glu Glu Gly His | | |
| 545 | 550 | 555 |
| Tyr Pro Pro Ala Pro Pro Pro Tyr Ser Glu Thr Asp Ser Gln Ala Ser | | |
| 565 | 570 | 575 |
| Arg Glu Arg Arg Met Lys Lys Asn Leu Ala Leu Ser Arg Glu Ser Leu | | |
| 580 | 585 | 590 |
| Val Val | | |

<210> 18
 <211> 575
 <212> PRT
 <213> Mus musculus

<400> 18
 Met Ala Pro Ala Ala Ser Ala Cys Ala Gly Ala Pro Gly Ser His Pro
 1 5 10 15
 Ala Thr Thr Ile Phe Val Cys Leu Phe Leu Ile Ile Tyr Cys Pro Asp
 20 25 30
 Arg Ala Ser Ala Ile Gln Val Thr Val Pro Asp Pro Tyr His Val Val
 35 40 45
 Ile Leu Phe Gln Pro Val Thr Leu His Cys Thr Tyr Gln Met Ser Asn
 50 55 60
 Thr Leu Thr Ala Pro Ile Val Ile Trp Lys Tyr Lys Ser Phe Cys Arg
 65 70 75 80
 Asp Arg Val Ala Asp Ala Phe Ser Pro Ala Ser Val Asp Asn Gln Leu
 85 90 95

Asn Ala Gln Leu Ala Ala Gly Asn Pro Gly Tyr Asn Pro Tyr Val Glu
 100 105 110
 Cys Gln Asp Ser Val Arg Thr Val Arg Val Val Ala Thr Lys Gln Gly
 115 120 125
 Asn Ala Val Thr Leu Gly Asp Tyr Tyr Gln Gly Arg Arg Ile Thr Ile
 130 135 140
 Thr Gly Asn Ala Gly Leu Thr Phe Glu Gln Thr Ala Trp Gly Asp Ser
 145 150 155 160
 Gly Val Tyr Tyr Cys Ser Val Val Ser Ala Gln Asp Leu Asp Gly Asn
 165 170 175
 Asn Glu Ala Tyr Ala Glu Leu Ile Val Leu Asp Trp Leu Phe Val Val
 180 185 190
 Val Val Cys Leu Ala Ser Leu Leu Phe Phe Leu Leu Leu Gly Ile Cys
 195 200 205
 Trp Cys Gln Cys Cys Pro His Thr Cys Cys Cys Tyr Val Arg Cys Pro
 210 215 220
 Cys Cys Pro Asp Lys Cys Cys Cys Pro Glu Ala Leu Tyr Ala Ala Gly
 225 230 235 240
 Lys Ala Ala Thr Ser Gly Val Pro Ser Ile Tyr Ala Pro Ser Ile Tyr
 245 250 255
 Thr His Leu Ser Pro Ala Lys Thr Pro Pro Pro Pro Ala Met Ile
 260 265 270
 Pro Met Arg Pro Pro Tyr Gly Tyr Pro Gly Asp Phe Asp Arg Thr Ser
 275 280 285
 Ser Val Gly Gly His Ser Ser Gln Val Pro Leu Leu Arg Glu Val Asp
 290 295 300
 Gly Ser Val Ser Ser Glu Val Arg Ser Gly Tyr Arg Ile Gln Ala Asn
 305 310 315 320
 Gln Gln Asp Asp Ser Met Arg Val Leu Tyr Tyr Met Glu Lys Glu Leu
 325 330 335
 Ala Asn Phe Asp Pro Ser Arg Pro Gly Pro Pro Asn Gly Arg Val Glu
 340 345 350
 Arg Ala Met Ser Glu Val Thr Ser Leu His Glu Asp Asp Trp Arg Ser
 355 360 365
 Arg Pro Ser Arg Ala Pro Ala Leu Thr Pro Ile Arg Asp Glu Glu Trp
 370 375 380
 Asn Arg His Ser Pro Arg Ser Pro Arg Thr Trp Glu Gln Glu Pro Leu
 385 390 395 400
 Gln Glu Gln Pro Arg Gly Gly Trp Gly Ser Gly Arg Pro Arg Ala Arg
 405 410 415
 Ser Val Asp Ala Leu Asp Asp Ile Asn Arg Pro Gly Ser Thr Glu Ser
 420 425 430
 Gly Arg Ser Ser Pro Pro Ser Ser Gly Arg Arg Gly Arg Ala Tyr Ala
 435 440 445
 Pro Pro Arg Ser Arg Ser Arg Asp Asp Leu Tyr Asp Pro Asp Asp Pro
 450 455 460
 Arg Asp Leu Pro His Ser Arg Asp Pro His Tyr Tyr Asp Asp Leu Arg
 465 470 475 480
 Ser Arg Asp Pro Arg Ala Asp Pro Arg Ser Arg Gln Arg Ser His Asp
 485 490 495
 Pro Arg Asp Ala Gly Phe Arg Ser Arg Asp Pro Gln Tyr Asp Gly Arg
 500 505 510
 Leu Leu Glu Glu Ala Leu Lys Lys Lys Gly Ala Gly Glu Arg Arg Arg
 515 520 525
 Val Tyr Arg Glu Glu Glu Glu Glu Glu Glu Gly His Tyr Pro Pro
 530 535 540
 Ala Pro Pro Pro Tyr Ser Glu Thr Asp Ser Gln Ala Ser Arg Glu Arg
 545 550 555 560
 Arg Met Lys Lys Asn Leu Ala Leu Ser Arg Glu Ser Leu Val Val

565

570

575

<210> 19
 <211> 526
 <212> PRT
 <213> Mus musculus

<400> 19
 Met Ala Pro Ala Ala Ser Ala Cys Ala Gly Ala Pro Gly Ser His Pro
 1 5 10 15
 Ala Thr Thr Ile Phe Val Cys Leu Phe Leu Ile Ile Tyr Cys Pro Asp
 20 25 30
 Arg Ala Ser Ala Ile Gln Val Thr Val Pro Asp Pro Tyr His Val Val
 35 40 45
 Ile Leu Phe Gln Pro Val Thr Leu His Cys Thr Tyr Gln Met Ser Asn
 50 55 60
 Thr Leu Thr Ala Pro Ile Val Ile Trp Lys Tyr Lys Ser Phe Cys Arg
 65 70 75 80
 Asp Arg Val Ala Asp Ala Phe Ser Pro Ala Ser Val Asp Asn Gln Leu
 85 90 95
 Asn Ala Gln Leu Ala Ala Gly Asn Pro Gly Tyr Asn Pro Tyr Val Glu
 100 105 110
 Cys Gln Asp Ser Val Arg Thr Val Arg Val Val Ala Thr Lys Gln Gly
 115 120 125
 Asn Ala Val Thr Leu Gly Asp Tyr Tyr Gln Gly Arg Arg Ile Thr Ile
 130 135 140
 Thr Gly Asn Ala Gly Leu Thr Phe Glu Gln Thr Ala Trp Gly Asp Ser
 145 150 155 160
 Gly Val Tyr Tyr Cys Ser Val Val Ser Ala Gln Asp Leu Asp Gly Asn
 165 170 175
 Asn Glu Ala Tyr Ala Glu Leu Ile Val Leu Val Tyr Ala Ala Gly Lys
 180 185 190
 Ala Ala Thr Ser Gly Val Pro Ser Ile Tyr Ala Pro Ser Ile Tyr Thr
 195 200 205
 His Leu Ser Pro Ala Lys Thr Pro Pro Pro Pro Pro Ala Met Ile Pro
 210 215 220
 Met Arg Pro Pro Tyr Gly Tyr Pro Gly Asp Phe Asp Arg Thr Ser Ser
 225 230 235 240
 Val Gly Gly His Ser Ser Gln Val Pro Leu Leu Arg Glu Val Asp Gly
 245 250 255
 Ser Val Ser Ser Glu Val Arg Ser Gly Tyr Arg Ile Gln Ala Asn Gln
 260 265 270
 Gln Asp Asp Ser Met Arg Val Leu Tyr Tyr Met Glu Lys Glu Leu Ala
 275 280 285
 Asn Phe Asp Pro Ser Arg Pro Gly Pro Pro Asn Gly Arg Val Glu Arg
 290 295 300
 Ala Met Ser Glu Val Thr Ser Leu His Glu Asp Asp Trp Arg Ser Arg
 305 310 315 320
 Pro Ser Arg Ala Pro Ala Leu Thr Pro Ile Arg Asp Glu Glu Trp Asn
 325 330 335
 Arg His Ser Pro Arg Ser Pro Arg Thr Trp Glu Gln Glu Pro Leu Gln
 340 345 350
 Glu Gln Pro Arg Gly Gly Trp Gly Ser Gly Arg Pro Arg Ala Arg Ser
 355 360 365
 Val Asp Ala Leu Asp Asp Ile Asn Arg Pro Gly Ser Thr Glu Ser Gly
 370 375 380
 Arg Ser Ser Pro Pro Ser Ser Gly Arg Arg Gly Arg Ala Tyr Ala Pro
 385 390 395 400
 Pro Arg Ser Arg Ser Arg Asp Asp Leu Tyr Asp Pro Asp Asp Pro Arg

<223> oligonucleotide anti sense primer

<400> 23
aggcggagat cgccagtcgt 20

<210> 24
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide sense primer

<400> 24
cctttgtcca cgtcgtttac gctc 24

<210> 25
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide anti sense primer

<400> 25
tcacagcggt gccctgcttg 20

<210> 26
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide sense primer

<400> 26
ttactgctcc gtggtctcag c 21

<210> 27
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide anti sense primer

<400> 27
agctactcct gtcaacgtct cc 22

<210> 28
<211> 167
<212> PRT
<213> Bos taurus

<400> 28
Met Arg Cys Gly Pro Leu Tyr Arg Phe Leu Trp Leu Trp Pro Tyr Leu
1 5 10 15
Ser Tyr Val Glu Ala Val Pro Ile Arg Lys Val Gln Asp Asp Thr Lys
20 25 30

Thr Leu Ile Lys Thr Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr
 35 40 45
 Gln Ser Val Ser Ser Lys Gln Arg Val Thr Gly Leu Asp Phe Ile Pro
 50 55 60
 Gly Leu His Pro Leu Leu Ser Leu Ser Lys Met Asp Gln Thr Leu Ala
 65 70 75 80
 Ile Tyr Gln Gln Ile Leu Thr Ser Leu Pro Ser Arg Asn Val Val Gln
 85 90 95
 Ile Ser Asn Asp Leu Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala
 100 105 110
 Ala Ser Lys Ser Cys Pro Leu Pro Gln Val Arg Ala Leu Glu Ser Leu
 115 120 125
 Glu Ser Leu Gly Val Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val
 130 135 140
 Val Ala Leu Ser Arg Leu Gln Gly Ser Leu Gln Asp Met Leu Arg Gln
 145 150 155 160
 Leu Asp Leu Ser Pro Gly Cys
 165

<210> 29
 <211> 146
 <212> PRT
 <213> Canis familiaris

<400> 29
 Val Pro Ile Arg Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr
 1 5 10 15
 Ile Val Ala Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser Ser
 20 25 30
 Lys Gln Arg Val Ala Gly Leu Asp Phe Ile Pro Gly Leu Gln Pro Val
 35 40 45
 Leu Ser Leu Ser Arg Met Asp Gln Thr Leu Ala Ile Tyr Gln Gln Ile
 50 55 60
 Leu Asn Ser Leu His Ser Arg Asn Val Val Gln Ile Ser Asn Asp Leu
 65 70 75 80
 Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala Ser Ser Lys Ser Cys
 85 90 95
 Pro Leu Pro Arg Ala Arg Gly Leu Glu Thr Phe Glu Ser Leu Gly Gly
 100 105 110
 Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val Val Ala Leu Ser Arg
 115 120 125
 Leu Gln Ala Ala Leu Gln Asp Met Leu Arg Arg Leu Asp Leu Ser Pro
 130 135 140
 Gly Cys
 145

<210> 30
 <211> 163
 <212> PRT
 <213> Gallus gallus

<400> 30
 Met Cys Trp Arg Pro Leu Cys Arg Leu Trp Ser Tyr Leu Val Tyr Val
 1 5 10 15
 Gln Ala Val Pro Cys Gln Ile Phe Gln Asp Asp Thr Lys Thr Leu Ile
 20 25 30
 Lys Thr Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr Ser Val Ser
 35 40 45
 Ala Lys Gln Arg Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro

50 55 60
 Ile Leu Ser Leu Ser Lys Met Asp Gln Thr Leu Ala Val Tyr Gln Gln
 65 70 75 80
 Val Leu Thr Ser Leu Pro Ser Gln Asn Val Leu Gln Ile Ala Asn Asp
 85 90 95
 Leu Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala Phe Ser Lys Ser
 100 105 110
 Cys Ser Leu Pro Gln Thr Ser Gly Leu Gln Lys Pro Glu Ser Leu Asp
 115 120 125
 Gly Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val Val Ala Leu Ser
 130 135 140
 Arg Leu Gln Gly Ser Leu Gln Asp Ile Leu Gln Gln Leu Asp Ile Ser
 145 150 155 160
 Pro Glu Cys

<210> 31
 <211> 146
 <212> PRT
 <213> Gorilla gorilla

<400> 31
 Val Pro Ile Gln Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr
 1 5 10 15
 Ile Val Thr Arg Ile Ser Asp Ile Ser His Thr Gln Ser Val Ser Ser
 20 25 30
 Lys Gln Lys Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro Ile
 35 40 45
 Leu Thr Leu Ser Lys Met Asp Gln Thr Leu Ala Val Tyr Gln Gln Ile
 50 55 60
 Leu Thr Ser Met Pro Ser Arg Asn Met Ile Gln Ile Ser Asn Asp Leu
 65 70 75 80
 Glu Asn Leu Arg Asp Leu Leu His Val Leu Ala Phe Ser Lys Ser Cys
 85 90 95
 His Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly Gly
 100 105 110
 Val Leu Glu Ala Ser Gly Tyr Ser Thr Glu Val Val Ala Leu Ser Arg
 115 120 125
 Leu Gln Gly Ser Leu Gln Asp Met Leu Trp Gln Leu Asp Leu Ser Pro
 130 135 140
 Gly Cys
 145

<210> 32
 <211> 167
 <212> PRT
 <213> Homo sapiens

<400> 32
 Met His Trp Gly Thr Leu Cys Gly Phe Leu Trp Leu Trp Pro Tyr Leu
 1 5 10 15
 Phe Tyr Val Gln Ala Val Pro Ile Gln Lys Val Gln Asp Asp Thr Lys
 20 25 30
 Thr Leu Ile Lys Thr Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr
 35 40 45
 Gln Ser Val Ser Ser Lys Gln Lys Val Thr Gly Leu Asp Phe Ile Pro
 50 55 60
 Gly Leu His Pro Ile Leu Thr Leu Ser Lys Met Asp Gln Thr Leu Ala
 65 70 75 80
 Val Tyr Gln Gln Ile Leu Thr Ser Met Pro Ser Arg Asn Val Ile Gln

Ile Ala Asn Asp Leu Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala
100 105 110
Phe Ser Lys Ser Cys Ser Leu Pro Gln Thr Ser Gly Leu Gln Lys Pro
115 120 125
Glu Ser Leu Asp Gly Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val
130 135 140
Val Ala Leu Ser Arg Leu Gln Gly Ser Leu Gln Asp Ile Leu Gln Gln
145 150 155 160
Leu Asp Val Ser Pro Glu Cys
165

<210> 35
<211> 146
<212> PRT
<213> Ovus aries

<400> 35
Val Pro Ile Arg Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr
1 5 10 15
Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser Ser
20 25 30
Lys Gln Arg Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro Leu
35 40 45
Leu Ser Leu Ser Lys Met Asp Gln Thr Leu Ala Ile Tyr Gln Gln Ile
50 55 60
Leu Ala Ser Leu Pro Ser Arg Asn Val Ile Gln Ile Ser Asn Asp Leu
65 70 75 80
Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala Ala Ser Lys Ser Cys
85 90 95
Pro Leu Pro Gln Val Arg Ala Leu Glu Ser Leu Glu Ser Leu Gly Val
100 105 110
Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val Val Ala Leu Ser Arg
115 120 125
Leu Gln Gly Ser Leu Gln Asp Met Leu Arg Gln Leu Asp Leu Ser Pro
130 135 140
Gly Cys
145

<210> 36
<211> 146
<212> PRT
<213> Pan troglodytes

<400> 36
Val Pro Ile Gln Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr
1 5 10 15
Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser Ser
20 25 30
Lys Gln Lys Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro Ile
35 40 45
Leu Thr Leu Ser Lys Met Asp Gln Thr Leu Ala Val Tyr Gln Gln Ile
50 55 60
Leu Thr Ser Met Pro Ser Arg Asn Met Ile Gln Ile Ser Asn Asp Leu
65 70 75 80
Glu Asn Leu Arg Asp Leu Leu His Val Leu Ala Phe Ser Lys Ser Cys
85 90 95
His Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly Gly
100 105 110
Val Leu Glu Ala Ser Gly Tyr Ser Thr Glu Val Val Ala Leu Ser Arg

115 120 125
 Leu Gln Gly Ser Leu Gln Asp Met Leu Trp Gln Leu Asp Leu Ser Pro
 130 135 140
 Gly Cys
 145

<210> 37
 <211> 146
 <212> PRT
 <213> Pongo pygmaeus

<400> 37
 Val Pro Ile Gln Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr
 1 5 10 15
 Val Ile Thr Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser Ser
 20 25 30
 Lys Gln Lys Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro Ile
 35 40 45
 Leu Thr Leu Ser Lys Met Asp Gln Thr Leu Ala Val Tyr Gln Gln Ile
 50 55 60
 Leu Thr Ser Met Pro Ser Arg Asn Val Ile Gln Ile Ser Asn Asp Leu
 65 70 75 80
 Glu Asn Leu Arg Asp Leu Leu His Val Leu Ala Phe Ser Lys Ser Cys
 85 90 95
 His Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu Asp Arg Leu Gly Gly
 100 105 110
 Val Leu Glu Ala Ser Gly Tyr Ser Thr Glu Val Val Ala Leu Ser Arg
 115 120 125
 Leu Gln Arg Ser Leu Gln Asp Met Leu Trp Gln Leu Asp Leu Ser Pro
 130 135 140
 Gly Cys
 145

<210> 38
 <211> 167
 <212> PRT
 <213> Rattus norvegicus

<400> 38
 Met Cys Trp Arg Pro Leu Cys Arg Phe Leu Trp Leu Trp Ser Tyr Leu
 1 5 10 15
 Ser Tyr Val Gln Ala Val Pro Ile His Lys Val Gln Asp Asp Thr Lys
 20 25 30
 Thr Leu Ile Lys Thr Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr
 35 40 45
 Gln Ser Val Ser Ala Arg Gln Arg Val Thr Gly Leu Asp Phe Ile Pro
 50 55 60
 Gly Leu His Pro Ile Leu Ser Leu Ser Lys Met Asp Gln Thr Leu Ala
 65 70 75 80
 Val Tyr Gln Gln Ile Leu Thr Ser Leu Pro Ser Gln Asn Val Leu Gln
 85 90 95
 Ile Ala His Asp Leu Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala
 100 105 110
 Phe Ser Lys Ser Cys Ser Leu Pro Gln Thr Arg Gly Leu Gln Lys Pro
 115 120 125
 Glu Ser Leu Asp Gly Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val
 130 135 140
 Val Ala Leu Ser Arg Leu Gln Gly Ser Leu Gln Asp Ile Leu Gln Gln
 145 150 155 160

Leu Asp Leu Ser Pro Glu Cys
165

<210> 39
<211> 167
<212> PRT
<213> Sus scrofa

<400> 39
Met Arg Cys Gly Pro Leu Cys Arg Phe Leu Trp Leu Trp Pro Tyr Leu
1 5 10 15
Ser Tyr Val Glu Ala Val Pro Ile Trp Arg Val Gln Asp Asp Thr Lys
20 25 30
Thr Leu Ile Lys Thr Ile Val Thr Arg Ile Ser Asp Ile Ser His Met
35 40 45
Gln Ser Val Ser Ser Lys Gln Arg Val Thr Gly Leu Asp Phe Ile Pro
50 55 60
Gly Leu His Pro Val Leu Ser Leu Ser Lys Met Asp Gln Thr Leu Ala
65 70 75 80
Ile Tyr Gln Gln Ile Leu Thr Ser Leu Pro Ser Arg Asn Val Ile Gln
85 90 95
Ile Ser Asn Asp Leu Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala
100 105 110
Ser Ser Lys Ser Cys Pro Leu Pro Gln Ala Arg Ala Leu Glu Thr Leu
115 120 125
Glu Ser Leu Gly Gly Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val
130 135 140
Val Ala Leu Ser Arg Leu Gln Gly Ala Leu Gln Asp Met Leu Arg Gln
145 150 155 160
Leu Asp Leu Ser Pro Gly Cys
165

<210> 40
<211> 4
<212> PRT
<213> Homo sapiens

<400> 40
Glu Thr Leu Asp
1

<210> 41
<211> 4
<212> PRT
<213> Mus musculus

<400> 41
Gln Lys Pro Glu
1

<210> 42
<211> 6
<212> PRT
<213> Homo sapiens

<400> 42
Leu Asp Ser Leu Gly Gly
1 5

<210> 43
<211> 4
<212> PRT
<213> Homo sapiens

<400> 43
Glu Lys Leu Glu
1

<210> 44
<211> 4
<212> PRT
<213> Homo sapiens

<400> 44
Glu Lys Pro Glu
1

<210> 45
<211> 4
<212> PRT
<213> Homo sapiens

<400> 45
Glu Lys Pro Asp
1

<210> 46
<211> 5
<212> PRT
<213> Homo sapiens

<400> 46
Thr Pro Asp Ser Leu
1 5

<210> 47
<211> 9
<212> PRT
<213> Homo sapiens

<400> 47
Gly Leu Gln Thr Leu Asp Ser Leu Gly
1 5

<210> 48
<211> 5
<212> PRT
<213> Homo sapiens

<400> 48
Gly Gly Val Leu Glu
1 5

<210> 49
<211> 6
<212> PRT
<213> Homo sapiens

<400> 49
Thr Pro Asp Ser Leu Gly
1 5

<210> 50
<211> 9
<212> PRT
<213> Homo sapiens

<400> 50
Ser Leu Gly Gly Val Leu Glu Ala Ser
1 5

<210> 51
<211> 6
<212> PRT
<213> Homo sapiens

<400> 51
Pro Glu Ser Leu Gly Gly
1 5

<210> 52
<211> 6
<212> PRT
<213> Homo sapiens

<400> 52
Pro Asp Ser Leu Gly Gly
1 5

<210> 53
<211> 7
<212> PRT
<213> Homo sapiens

<400> 53
Leu Gly Gly Val Leu Glu Ala
1 5

<210> 54
<211> 22
<212> PRT
<213> Homo sapiens

<400> 54
Glu Asn Leu Arg Asp Leu Leu His Val Leu Ala Phe Ser Lys Ser Cys
1 5 10 15
His Leu Pro Trp Ala Ser
20

<210> 55
<211> 22
<212> PRT
<213> Homo sapiens

<400> 55
Leu Leu His Val Leu Ala Phe Ser Lys Ser Cys His Leu Pro Trp Ala
1 5 10 15

Ser Gly Leu Glu Thr Leu
20

<210> 56
<211> 22
<212> PRT
<213> Homo sapiens

<400> 56
Ala Phe Ser Lys Ser Cys His Leu Pro Trp Ala Ser Gly Leu Glu Thr
1 5 10 15
Leu Asp Ser Leu Gly Gly
20

<210> 57
<211> 22
<212> PRT
<213> Homo sapiens

<400> 57
Cys His Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly
1 5 10 15
Gly Val Leu Glu Ala Ser
20

<210> 58
<211> 18
<212> PRT
<213> Homo sapiens

<400> 58
Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly Gly Val
1 5 10 15
Leu Glu

<210> 59
<211> 14
<212> PRT
<213> Homo sapiens

<400> 59
Trp Ala Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly Gly Val
1 5 10

<210> 60
<211> 21
<212> PRT
<213> Homo sapiens

<400> 60
Ala Ser Gly Leu Glu Thr Asp Ser Leu Gly Gly Val Leu Glu Ala Ser
1 5 10 15
Gly Tyr Ser Thr Glu
20

<210> 61
<211> 10
<212> PRT
<213> Homo sapiens

<400> 61
 Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly
 1 5 10

<210> 62
 <211> 22
 <212> PRT
 <213> Homo sapiens

<400> 62
 Thr Leu Asp Ser Leu Gly Gly Val Leu Glu Ala Ser Gly Tyr Ser Thr
 1 5 10 15
 Glu Val Val Ala Leu Ser
 20

<210> 63
 <211> 22
 <212> PRT
 <213> Homo sapiens

<400> 63
 Gly Gly Val Leu Glu Ala Ser Gly Tyr Ser Thr Glu Val Val Ala Leu
 1 5 10 15
 Ser Arg Gly Gln Gly Ser
 20

<210> 64
 <211> 22
 <212> PRT
 <213> Mus musculus

<400> 64
 Glu Asn Leu Arg Asp Leu Leu His Leu Leu Ala Phe Ser Lys Ser Cys
 1 5 10 15
 Ser Leu Pro Gln Thr Ser
 20

<210> 65
 <211> 22
 <212> PRT
 <213> Mus musculus

<400> 65
 Leu Leu His Leu Leu Ala Phe Ser Lys Ser Cys Ser Leu Pro Gln Thr
 1 5 10 15
 Ser Gly Leu Gln Lys Pro
 20

<210> 66
 <211> 22
 <212> PRT
 <213> Mus musculus

<400> 66
 Ala Phe Ser Lys Ser Cys Ser Leu Pro Gln Thr Ser Gly Leu Gln Lys
 1 5 10 15
 Pro Glu Ser Leu Asp Gly
 20

<210> 67
<211> 22
<212> PRT
<213> Mus musculus

<400> 67
Cys Ser Leu Pro Gln Thr Ser Gly Leu Gln Lys Pro Glu Ser Leu Asp
1 5 10 15
Gly Val Leu Glu Ala Ser
20

<210> 68
<211> 18
<212> PRT
<213> Mus musculus

<400> 68
Leu Pro Gln Thr Ser Gly Leu Gln Lys Pro Glu Ser Leu Asp Gly Val
1 5 10 15
Leu Glu

<210> 69
<211> 14
<212> PRT
<213> Mus musculus

<400> 69
Gln Thr Ser Gly Leu Gln Lys Pro Glu Ser Leu Asp Gly Val
1 5 10

<210> 70
<211> 22
<212> PRT
<213> Mus musculus

<400> 70
Thr Ser Gly Leu Gln Lys Pro Glu Ser Leu Asp Gly Val Leu Glu Ala
1 5 10 15
Ser Leu Tyr Ser Thr Glu
20

<210> 71
<211> 10
<212> PRT
<213> Mus musculus

<400> 71
Ser Gly Leu Gln Lys Pro Glu Ser Leu Asp
1 5 10

<210> 72
<211> 22
<212> PRT
<213> Mus musculus

<400> 72
Lys Pro Glu Ser Leu Asp Gly Val Leu Glu Ala Ser Leu Tyr Ser Thr
1 5 10 15

Glu Val Val Ala Leu Ser
20

<210> 73
<211> 22
<212> PRT
<213> Mus musculus

<400> 73
Asp Gly Val Leu Glu Ala Ser Leu Tyr Ser Thr Glu Val Val Ala Leu
1 5 10 15
Ser Arg Leu Gln Gly Ser
20

<210> 74
<211> 67
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide Chimeric oligonucleotides

<400> 74
atgcaacagg acggacttgg agtagttttc uacuccaagt cagtccuguu gcaugcgcgt 60
ttcgcgc 67

<210> 75
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide Forward Primer

<400> 75
tgtccacgtc gtttacgctc 20

<210> 76
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide Reverse Primer

<400> 76
tcccacttcc gttccttgctc 20

<210> 77
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> oligonucleotide Probes endogenous/mutant

<400> 77
cctactccaa gtcmgctcctg ttgcatt 27

<210> 78
 <211> 67
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Chimeric oligonucleotides

 <400> 78
 gaccctgccc tgtacctacc taccagatgt tttcaucugg uagggttcagg gcagggucgc 60
 gcggtttt 67

 <210> 79
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Forward Primer

 <400> 79
 gtggtgatcc tcttccagcc t 21

 <210> 80
 <211> 19
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Reverse Primer

 <400> 80
 ccagatgacg atggggttgc 19

 <210> 81
 <211> 25
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Probes endogenous/mutant

 <400> 81
 accctgccct gwcctaccag atgac 25

 <210> 82
 <211> 68
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Chimeric oligonucleotides

 <400> 82
 tggctgagct cttacctggt tttcattttt gaaaaccagg tcagagctca gccagcgcg 60
 tttcgcgc 68

 <210> 83
 <211> 20

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Forward Primer

 <400> 83
 gagctcatcg tccttgggag 20

 <210> 84
 <211> 19
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Reverse Primer

 <400> 84
 agtcttctat gggccccgc 19

 <210> 85
 <211> 27
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Probes endogenous/mutant

 <400> 85
 caccgactcg agamtggacc aaaagtc 27

 <210> 86
 <211> 68
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Chimeric oligonucleotides

 <400> 86
 ggttggtgta tgcctggctg ccttcttttg aaggcagcca gtcataccac aaccgcgcgt 60
 tttcgcgc 68

 <210> 87
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Forward Primer

 <400> 87
 acgcagagct catcgctcctt 20

 <210> 88
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> oligonucleotide Reverse Primer

 <400> 88
 gatgcccagg aggaggaaga 20

 <210> 89
 <211> 23
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Probes endogenous/mutant

 <400> 89
 caacaccata ckgaccgacg gaa 23

 <210> 90
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide mouse LSR specific primer

 <400> 90
 acgcatggga atcatggc 18

 <210> 91
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <400> 91
 taggggtgag cggcgggg 18

 <210> 92
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <220>
 <221> misc_feature
 <222> 10..12
 <223> n=a, g, c or t

 <400> 92
 gagggctggn nntaggggtg a 21

 <210> 93
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <220>
 <221> misc_feature
 <222> 10..11
 <223> n=a, g, c or t

 <400> 93
 agggctgggn ntaggggtga 20

 <210> 94
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <400> 94
 gtgggagccg agggctgg 18

 <210> 95
 <211> 19
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <220>
 <221> misc_feature
 <222> 10
 <223> n=a, g, c or t

 <400> 95
 gtgggagccn agggctggg 19

 <210> 96
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <400> 96
 gcggcggccg ggtgggag 18

 <210> 97
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

 <400> 97

ttggccggag cagatggg 18

<210> 98
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

<220>
 <221> misc_feature
 <222> 10..11
 <223> n=a, g, c or t

<400> 98
 gcagatgggn nccggaaggg 20

<210> 99
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

<220>
 <221> misc_feature
 <222> 10..12
 <223> n=a, g, c or t

<400> 99
 agggctgggn nnaggggtga g 21

<210> 100
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

<220>
 <221> misc_feature
 <222> 10..12
 <223> n=a, g, c or t

<400> 100
 aggggtgagn nncggggagg g 21

<210> 101
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> oligonucleotide Zinc finger nuclotides of SEQID1

<400> 101

| | |
|---|----|
| aagtgggtct cggttgca | 18 |
| <210> 102 | |
| <211> 21 | |
| <212> DNA | |
| <213> Artificial Sequence | |
| <220> | |
| <223> oligonucleotide zinc finger LSR sequences | |
| <400> 102 | |
| aaggtcgctt atggtgcaga c | 21 |
| <210> 103 | |
| <211> 20 | |
| <212> DNA | |
| <213> Artificial Sequence | |
| <220> | |
| <223> oligonucleotide zinc finger LSR sequences | |
| <400> 103 | |
| gtgggagccc gggggctgga | 20 |
| <210> 104 | |
| <211> 18 | |
| <212> DNA | |
| <213> Artificial Sequence | |
| <220> | |
| <223> oligonucleotide zinc finger LSR sequences | |
| <400> 104 | |
| tgggggtggg cggcgggg | 18 |
| <210> 105 | |
| <211> 20 | |
| <212> DNA | |
| <213> Artificial Sequence | |
| <220> | |
| <223> oligonucleotide zinc finger LSR sequences | |
| <400> 105 | |
| ccgggagtgc gcagggggta | 20 |
| <210> 106 | |
| <211> 19 | |
| <212> DNA | |
| <213> Artificial Sequence | |
| <220> | |
| <223> oligonucleotide zinc finger LSR sequences | |
| <400> 106 | |
| gtggctgcac aaggtcgcc | 19 |